



**Helder Duarte Paixão
Berenguer**

**Predisposição de *Eucalyptus* a *Neofusicoccum
kwambonambiense* causada por stress hídrico**

***Eucalyptus* predisposition to *Neofusicoccum
kwambonambiense* under water stress**

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Dissertação apresentada à Universidade de Aveiro para cumprimento dos requisitos necessários à obtenção do grau de Mestre em Biologia Molecular e Celular, realizada sob a orientação científica do Doutor Artur Jorge da Costa Peixoto Alves, investigador principal do Departamento de Biologia da Universidade de Aveiro e co-orientação da Doutora Glória Catarina Cintra da Costa Pinto, professora auxiliar convidada do Departamento de Biologia da Universidade de Aveiro.

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To curiosity, the spark behind science

o júri

presidente

Prof. Doutora Maria Paula Polónia Gonçalves
Professora Associada do Departamento de Biologia da Universidade de Aveiro

arguente principal

Doutora Lia-Tânia Rosa Dinis
Investigadora de Pós-Doutoramento da Universidade de Trás-os-Montes e Alto Douro

orientador

Prof. Doutor Artur Jorge da Costa Peixoto Alves
Investigador Principal do Departamento de Biologia e CESAM da Universidade de Aveiro

agradecimentos

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palavras-chave

Floresta, Desempenho das plantas, *Eucalyptus globulus*, Interações bióticas e abióticas, Interações planta-patogénio

resumo

Em Portugal, o eucalipto, particularmente o *Eucalyptus globulus*, ocupa mais de 800 000 ha. Devido a ser uma importante fonte de biomassa para painéis de fibras, carvão industrial, lenha e pasta de papel, tornou-se um género chave de considerável importância económica. Contudo, a produtividade de *E. globulus* tem encontrado novas pressões, sendo a seca resultante das alterações climáticas, uma das mais hostis. A seca pode levar a uma diminuição do crescimento e produtividade: diretamente; ou indiretamente através do aumento da suscetibilidade a agentes patogénicos através da predisposição. O fungo ascomiceto *Neofusicoccum kwambonambiense* é um agente fitopatogénico endofítico oportunista que se sabe afetar severamente *E. globulus*, e cuja presença já fora confirmada em Portugal. Tomando tal em consideração, torna-se importante avaliar o efeito de predisposição que a seca poderá ter na interação *N. kwambonambiense* - *E. globulus*. Para tal foram criados quatro grupos de tratamento: *E. globulus* foram primeiramente sujeitos a um período de aclimatização de 66 dias no qual foram periodicamente irrigados (80% de capacidade de campo). Seguidamente, dois grupos foram sujeitos a uma diminuição progressiva da irrigação. Os outros dois grupos permaneceram bem regados. Uma vez que os tratamentos stressados atingiram 18% de capacidade de campo (23 dias), um grupo bem regado e um grupo stressado foram inoculados com *N. kwambonambiense*. Todas os tratamentos foram mantidos nestas condições durante um período de 66 dias, findo o qual foi obtido um conjunto de parâmetros morfológicos, fisiológicos e bioquímicos. As plantas bem regadas, apesar de terem sido inoculadas com *N. kwambonambiense* apresentaram um aumento dos parâmetros fotossintéticos o que terá permitido a defesa da planta através de uma produção amplificada de açúcares, prolina e ácido salicílico. Danos oxidativos (parcialmente observados através do conteúdo em *malondialdeído*) foram evitados, em parte, devido à ação da prolina e açúcares solúveis. O *stress* hídrico levou a uma diminuição do crescimento confirmado pela redução do conteúdo em ácido-indole-acético. Ocorreu uma diminuição do potencial hídrico, a qual, em conjunto com o aumento do ácido abscísico, levou ao fecho dos estomas e diminuição da fotossíntese. Os danos oxidativos não foram controlados, afetando o estado do *E. globulus*. Ademais, o *N. kwambonambiense* provocou um aumento do conteúdo em ácido jasmónico, típico de agentes patogénicos necrotróficos, o que poderá sugerir que o fungo passou de um estilo de vida hemibiotrófico para necrotrófico, à medida que as células degeneravam. Os *E. globulus* stressados apresentavam maiores lesões externas e cancrios, conjuntamente com uma maior progressão interna do fungo. Os nossos resultados comprovam que a seca criou um melhor substrato para o desenvolvimento do fungo e diminuiu a capacidade de resposta da planta. Tal resultou num aumento da suscetibilidade e severidade da doença confirmando a predisposição.

keywords

Forest trees, Plant performance, *Eucalyptus globulus*, Biotic and abiotic interactions, Plant-pathogen interactions

abstract

In Portugal, *Eucalyptus*, particularly *Eucalyptus globulus*, occupies more than 800 000 ha and, due to being a major source of biomass for fiberboard, industrial charcoal, fuel wood and paper pulp, has become a key genus, with a considerable economic importance. However, *E. globulus* productivity faces new pressures, with climate change-driven drought as one of the most hostile ones. Drought can lead to growth impairment and yield reduction: directly; or indirectly, through the increase of plant susceptibility to pathogens by a predisposition mechanism. *Neofusicoccum kwambonambiense* is an endophytic opportunist phytopathogen known to severely affect *E. globulus*, whose incidence has already been reported in Portugal. Taking all in consideration, it is of major importance to assess the predisposition effect that drought may have on the *N. kwambonambiense* - *E. globulus* interaction. For such purpose, four treatment groups were established: *E. globulus* were firstly subjected to a 66-days acclimation period in which plants were periodically watered (80% of field capacity). After that, two groups were exposed to a progressive water supply restriction. The other two remained well-watered. Once water-stressed plants achieved 18% of field capacity (23 days), a well-watered and a water-stress group were inoculated with *N. kwambonambiense*. All treatments were kept in these conditions throughout a 65 days' period, at which moment a set of morphological, physiological and biochemical parameters was obtained. Well-watered plants, despite being infected with *N. kwambonambiense*, presented an overall photosynthetic increase, which enabled plant defense through the production of sugars, proline and salicylic acid. Oxidative damages (partially observed through malondialdehyde content), were avoided in part due to proline and soluble sugars. Water stress lead to a direct growth impairment confirmed through an indole-acetic-acid content decrease. A water-potential reduction occurred, which, together with abscisic acid, lead to stomatal closure and overall photosynthetic efficiency decline. Oxidative damages weren't properly managed and further affected *E. globulus*. Furthermore, *N. kwambonambiense* was found to promote a jasmonic acid content increase, typical of necrotrophic pathogens, which may suggest a lifestyle change from hemibiotrophic to necrotrophic as plant cells progressively degenerate. Ultimately, water-stressed *E. globulus* presented larger external lesion extensions and steam cankers and a superior internal fungi progression. Our results conclusively demonstrate that water stress created a better substrate for fungi development and decreased the plant's ability to respond. Such resulted in higher susceptibility and disease severity confirming predisposition.

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Part I

Forest and climate change

Forests supply energy, building materials, and food, but also deliver services such as carbon storing, biodiversity hosting, and climate regulation (Trumbore et al., 2015). Although adapted to some level of disturbance, forests face new threats, with human-related intensification of stressors, such as climate change, as one of the most hostile ones (Thuiller et al., 2005; Trumbore et al., 2015).

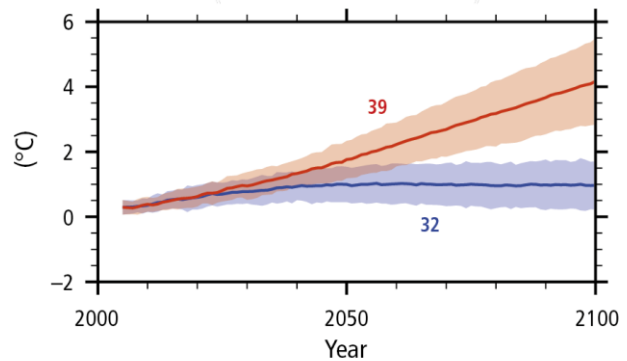


Figure 1 Global average surface temperature change from 2006 to 2100 determined by multi-model simulations under two scenarios RCP2.6 (blue) and RCP8.5 (red). Source: Pachauri et al., (2014)

From 1979 to 2010, global temperatures increased by about 0.5°C (Foster and Rahmstorf, 2011) and, according to the Intergovernmental Panel on Climate Change (IPCC) 2014 report, Earth's surface temperature is projected to continue to rise over the course of the 21st century (Pachauri et al., 2014) (fig.1), with more frequent temperature extremes and reductions in summer precipitation across Europe (Blenkinsop and Fowler, 2007; Dai et al., 2004). This combination will give rise to an increase of drought episodes, a major abiotic influence that greatly limits plant growth and development (Reddy et al., 2004). Additionally, climatic models predict that the Mediterranean area will be one of most affected regions (Giorgi, 2006), particularly, the northern Mediterranean, Lusitanian, and Mediterranean mountain regions (Thuiller et al., 2005).

Climate is an important distribution driver, predominantly regarding plant pathogenic fungi, being capable to give rise to new pathosystems (Desprez-Loustau et al., 2007; La Porta et al., 2008). It can exacerbate forest pathogens invasions, increasing the range of both native and exotic species (Jactel et al., 2012). This way, mild diseases, under normal conditions, can become severe under extreme weather conditions (Desprez-Loustau et al., 2006; Slippers and Wingfield, 2007). Such interaction gives rise to new and complex challenges to the eucalypt forest industry that are further discussed throughout this dissertation.

***Eucalyptus* spp. plantations**

Despite constituting only 7% of the global forested area, planted forests significantly contribute to the overall forest's goods and services (FAO, 2015). *Eucalyptus* spp. belong to the *Myrtaceae* family, are native to Australasia, and comprise more than 700 species (Brooker and Hopper, 2002). *Eucalyptus* spp. are known for their rapid growth, short rotation time, and high adaptability to soils, climate, and nutrient conditions (Booth, 2013; Florence, 2004). For such reasons, are considered a major source of biomass for paper pulp production, being used for industrial charcoal and firewood production, and recognized as a potential source of biofuels and biomaterials (Luger, 2003; Turnbull, 1999). Therefore, it comes with no surprise that *Eucalyptus* spp. are the most widely planted hardwood, explored in plantations in over 90 countries, extending from southern Europe to South Africa, totaling more than 20 million hectares (Flynn, 2010; Iglesias-Trabado and Wilstermann, 2008).

Eucalyptus globulus

One of the most noteworthy *Eucalyptus* spp. species is *Eucalyptus globulus* (Labill.). In Europe, it covers 1.3 million ha, mainly in the Iberian Peninsula (more than 80%), France and Italy (Iglesias-Trabado and Wilstermann, 2008). In spite of the non-optimal conditions of the Mediterranean climate, it has become the most widely introduced species in the region (Navarrete-Campos et al., 2013).

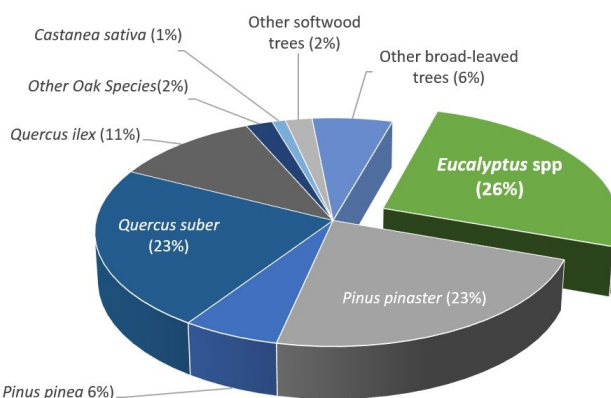


Figure 2 Forest area distribution by species/genera/group in mainland Portugal. Adapted from: ICNF (2013)

Eucalyptus globulus was introduced in Portugal around 1860, for timber and paper pulp production. Its expansion arose later, in the mid-20th century, along with the paper and pulp industry growth (Borrvalho et al., 2007). The combination of breeding programs and improved techniques for cutting rooting led to the selection of clones taking into account their high growth rates, high pulp yield and environmental adaptability (Costa et al., 2004). Since its introduction, *E. globulus*' planted area has progressively increased (fig. 2), reaching 811,943 ha in 2010 (ICNF, 2013), establishing *E. globulus* as the dominant forest type in Portugal (ICNF, 2013). *E. globulus*, due to the pulp industry, represents a valuable contribution to the country's economy (CELPA, 2014). Although, as an exotic species, *E. globulus* has been free from most diseases, this situation has changed in the last 30 years (Bragança et al., 2015).

The family *Botryosphaeriaceae*

Members of the *Botryosphaeriaceae* (*Botryosphaeriales*, Ascomycota) fungal family were first described in the 1820's and include more than 2000 taxa (Slippers and Wingfield, 2007). Presenting a worldwide distribution, they occur on a wide range of monocotyledonous, dicotyledonous and gymnosperms hosts (Slippers and Wingfield, 2007) and can infect healthy plants, directly through lenticels, stomata or other openings, and be present in a latent state (Michailides, 1991; Slippers and Wingfield, 2007). Infections of woody hosts are believed to predominantly occur through horizontal transmission via spores (Slippers and Wingfield, 2007). Considered opportunistic host colonists, they hold the ability to live endophytically in their hosts, becoming pathogenic and triggering disease when the host is exposed to a predisposing stress condition, such as drought, which is the most frequently reported one (Mohali et al., 2007; Slippers and Wingfield, 2007; Van Niekerk et al., 2011). *Botryosphaeriaceae* are commonly linked to cankers and dieback, followed by broad kino production, and, in severe circumstances, tree death (fig. 3) (Pavlic et al., 2007).



Figure 3 Typical *Neofusicoccum* and *Fusicoccum* symptoms on *Eucalyptus* spp. (a) and (b): tree death; (c) and (d): internal damage. Source: Slippers and Wingfield, (2007)

Their high occurrence, dispersion aptitude, and ability to cause disease when their hosts are under stress, make these pathogens a significant threat to forests and plantations, particularly to non-native species (Crous et al., 2006; Slippers and Wingfield, 2007). In Portugal, according to Barradas et al. (2016), nine species from three different *Botryosphaeriaceae* genera were found in *E. globulus*, namely *Botryosphaeria dothidea*, *Diplodia corticola*, *Diplodia seriata*, *Neofusicoccum* spp., *Neofusicoccum australe*, *Neofusicoccum algeriense*, *Neofusicoccum eucalyptorum*, *Neofusicoccum parvum*, and *Neofusicoccum kwambonambiense*.

Neofusicoccum kwambonambiense

Neofusicoccum species affect several woody species worldwide (Sakalidis et al., 2013). *Neofusicoccum kwambonambiense* (Pavlic, Slippers & M.J. Wingf.) is part of the *Neofusicoccum parvum* – *Neofusicoccum ribis* species complex, and is commonly found in its asexual stage (Pavlic et al., 2009).

Neofusicoccum kwambonambiense has been found on four continents, six countries and fourteen hosts (Sakalidis et al., 2013). Those including blueberry, in Florida (Wright and Harmon, 2010), *Syzygium cordatum* (Pavlic et al., 2007), *Cordia Africana* (Slippers et al., 2014), *Eucalyptus grandis* (Pillay et al., 2013) and *Acacia karro* (Slippers et al., 2009), in South Africa. *Eucalyptus dunnii* and *Corymbia torelliana*, in Australia (Sakalidis et al., 2011) and *Vitis vinifera*, in Uruguay (Abreo et al., 2013). From the nine *Botryosphaeriaceae* species detected in Portugal, in 2012, by Barradas et al. (2016), *N. kwambonambiense*, found causing stem canker (fig. 4), was considered to be the most aggressive. For such reason, *N. kwambonambiense* was the chosen pathogen to explore in this study.

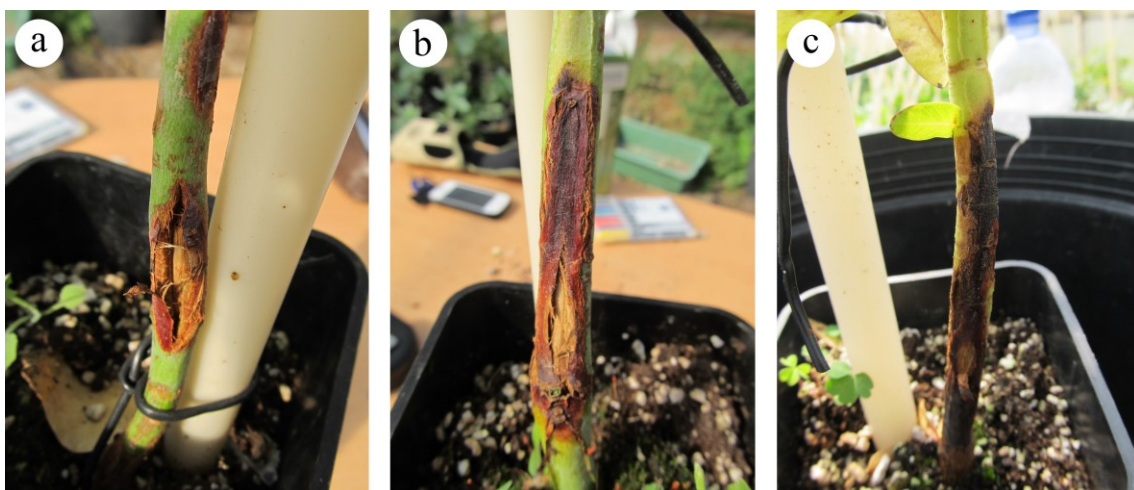


Figure 4 External lesion caused by *N. kwambonambiense* on *E. globulus* x *E. cypellocarpa* (a), *E. globulus* (b) and *E. nitens* (c) under a pot experiment. © Carla Barradas, 2014.

Quarantine systems have proven unsuccessful to control latent pathogens since they are able to be endophytically present in healthy plant material (Slippers and Wingfield, 2007), and are difficult to control once introduced into new areas (Swart and Wingfield, 1991). Chemical control is problematic on a large scale and removal or treatment of diseased tree parts, along with sanitation intended to reduce spore loads, appears to be the best approach to reduce disease (Brown-Rytlewski and McManus, 2000; Slippers and Wingfield, 2007; Stanosz et al., 2005). Nevertheless, symptoms are likely to continue to appear from other endophytic infections, and susceptible species can be replaced with resistant ones (Swart and Wingfield, 1991).

Drought-disease interaction effect on plant physiology

Plant predisposition and multiple stress concept

Interaction effects through host physiology can be allocated into two types of interaction: (1) combined effects of infection and drought on tree physiology: the “multiple stress concept”; and (2) an effect of drought on tree susceptibility to disease: the “predisposition concept” (Desprez-Loustau et al., 2006). Having in account that *N. kwambonambiense* is known to cause serious damage on plants previously subjected to drought, this study will focus on the predisposition effect.

The concept of predisposition, that is, the increased susceptibility to pathogen infection caused by environmental stress, was introduced and developed by Schoeneweiss (1978). Two main mechanisms may explain predisposition: (1) water stressed plant tissues are a better substrate for pathogen growth and (2): a decrease in plant active resistance may occur (Desprez-Loustau et al., 2006). This subject is of great scientific interest and has been reviewed (Bostock et al., 2014; Boyer, 1995; Desprez-Loustau et al., 2006). The great majority of published studies dealt with *Quercus*, *Populus* and *Pinus*, genus of considerable productive and environmental importance (Desprez-Loustau et al., 2006) and, conclusively, reported cases of water stress-caused predisposition. The hybrid *Populus nigra* × *Prunus maximowiczii* developed larger cankers as a result of *Septoria musiva* infection when grown under moderate drought (Franceschini et al., 2004); *Pinus resinosa*, under similar conditions, became more susceptible to *Sphaeropsis sapinea* infection (Blodgett and Stanosz, 1997); pistachio developed more severe *Botryosphaeria* blight disease (Ma et al., 2001); and, in field-grown grapevines, symptom development associated with *Botryosphaeriaceae*, was more severe in cases where plants have been, or still were, exposed to water-stress (Van Niekerk et al., 2011).

Plant response to drought and pathogen

Several investigations revealed that abiotic and biotic stressors can engage common signals and share responsive genes and products (Bostock et al., 2014). In recent years, research has largely focused on individual biotic or abiotic stress (Niinemets, 2010). Several studies regarding the effect of water stress on *Eucalyptus* spp. have been explored (Correia et al., 2016, 2014b; Li et al., 2000; Martorell et al., 2014; White et al., 1996a). Additionally, the distribution and effect of *Botryosphaeriaceae* on *Eucalyptus* spp. has also been addressed (Alves et al., 2013; Barradas et al., 2016; Pavlic et al., 2007; Pérez et al., 2010; Slippers and Wingfield, 2007). Nonetheless, the response to the collective drought and pathogen stress cannot be predicted based in the plant's response to the individual stresses, being guaranteed to lead to a more complex scenario (Desprez-Loustau et al., 2006; Niinemets, 2010; Rejeb et

al., 2014). The link between these three elements is a clear example of a “disease triangle” (fig. 5). Introduced by Horsfall and Dimond (1960), the “disease triangle” is a plant pathology concept that studies the interactions between a biotic agent (*N. kwambonambiense*), a susceptible host (*E. globulus*), and a favorable environment for disease development (climate change driven-drought). This work intends to fill the knowledge gap that exists regarding this scenario from a predisposition point of view and, although in an interaction study all factors response should be explored, the focus of the study was narrowed to the *E. globulus* physiological response.

Biomass, growth and morphology

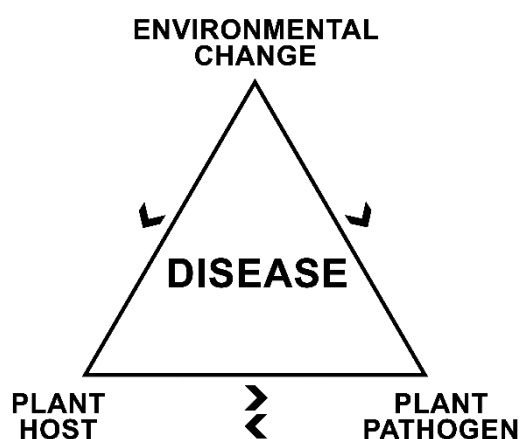


Figure 5 Modification of the environment-host-pathogen triangle highlighting the effect of environmental change on the interaction between plant host and plant pathogen. Adapted from: Grulke,(2011)

Plant growth and development impairment, as a result of turgor pressure reduction and decreased CO₂ diffusion, is one of the most drought sensitive and perceptible effects (Shao et al., 2008). Such causes water-stressed plants to often be smaller when compared to optimal watered ones (Souza and Cardoso, 2003), as found in *E. globulus* and other *Eucalyptus* spp. (Correia et al., 2014b; Costa et al., 2004; Li et al., 2000; Pita and Pardos, 2001; Shvaleva et al., 2006). Pathogens can also disturb both primary and secondary metabolism, affecting plant growth and development (Berger et al., 2007). Exposure to biotic and abiotic stresses thus leads to a disruption of the plant's metabolism, with physiological costs and, ultimately, to reductions in plant fitness and productivity (Rejeb et al., 2014).

Water relations and osmolyte accumulation

In order for the plant to absorb water, its water potential must be lower than that of the water supply. A reduction of the plant's water potential tends to occur under water deficit conditions (Boyer, 1982), with several *E. globulus* studies reporting such a response (Correia et al., 2014b; Pita et al., 2003; Shvaleva et al., 2006; White et al., 1996b). Although water

content may have a direct effect on fungal growth within the host, most fungal pathogens display plasticity and can develop at water potentials well below the minimum for growth of most higher plants (Desprez-Loustau et al., 2006). For example, *Botryosphaeria dothidea* exhibited a remarkable tolerance to low water potentials (Ma et al., 2001). In another work with *B. dothidea*, the author found that hyphae in vessels of stressed *Betula alba* stems were rectilinear and comparatively larger than the ones of unstressed plants (McPartland, 1984). Woody tissue desiccation leads to mechanical strength loss and can result in bark cracks that ease opportunistic endophytic pathogen invasion (Bettucci et al., 1999; Moore, 1987).

Osmotic adjustment is a physiological adaptation intended to maintain cell turgor and facilitate soil water extraction (Chen and Jiang, 2010; Morgan, 1984). It depends upon the accumulation of compatible solutes, particularly soluble sugars (Chen and Jiang, 2010; Couée et al., 2006). There is a strong correlation between water potential, soluble sugars concentration and stress tolerance. Drought stress leads to a water potential decrease and a soluble sugars accumulation (Morkunas and Ratajczak, 2014; Sperdouli and Moustakas, 2012). *E. globulus* showed an increase in sucrose upon water stress (Merchant et al., 2006). Additionally, soluble sugars have been found to express a signaling function, similar to hormones, upon pathogen interaction, regarding the induction of pattern-triggered immunity (PTI) and effector-triggered immunity (ETI) (Horacio and Martinez-Noel, 2013; Rojas et al., 2014; Rolland et al., 2006). Sugars are closely related to the abscisic acid signaling cascade, with such association leading to signal amplification (Pinheiro and Chaves, 2011). Thus, an effective immune response depends of a synchronized interaction of hormonal and sugar pathways that, under drought, may be severely impaired (Bolouri and Van den Ende, 2012). Proline is an amino acid essential for plant primary metabolism (Szabados and Savoure, 2010). This osmolyte is produced in response to stress, particularly drought (Choudhary et al., 2005; Sperdouli and Moustakas, 2012), protecting plant cells by osmotic adjustment and membrane stabilization (Sperdouli and Moustakas, 2012). Shvaleva et al. (2006) found an increase in proline concentration in water-stressed *E. globulus*. Proline has also been attributed an antioxidant role, suggesting a ROS scavenging activity (Matysik et al., 2002; Smirnov and Cumbes, 1989).

Photosynthetic performance and related pigments

Photosynthesis is amongst the primary processes to be affected by drought (Xu et al., 2010). Such may occur as a result of two mechanisms: (1) directly, through a CO₂ availability decrease, as a consequence of stomatal closure (Chaves et al., 2009) and metabolic impairment (Reddy et al., 2004); and (2), indirectly, and typical of multiple stress conditions,

as a result of secondary effects, specifically oxidative stress (Chaves et al., 2009; Chaves and Oliveira, 2004). In addition to the CO₂ diffusion reduction through the stomata, drought leads to a CO₂ mesophyll conductance and transpiration decline (Flexas et al., 2004; Lima et al., 2003). Guarnaschelli et al. (2006) showed a stomatal conductance decline on water-stressed *E. globulus* seedlings. Lima et al. (2003) had similar results in several *Eucalyptus* species. Photosynthesis efficiency decline causes a reduction in ATP production. That results in a decreased synthesis of enzymes and proteins that are involved in pathogen defense (e.g. enzymes with a lytic action on fungal hyphae and phytoalexins). Such ultimately leads to an increased pathogen susceptibility and disease severity (Desprez-Loustau et al., 2006; Kruger and Manion, 1994; Van Niekerk et al., 2011).

Chlorophylls content is positively associated with photosynthetic rate (Anjum et al., 2011). Drought is known to lead to a loss of chlorophyll content, as the plant is forced to divert the absorbed light to other processes, like thermal dissipation to protect the photosynthetic apparatus (Reddy et al., 2004). Nevertheless, such reaction is not identical in all plants. When *Picea abies* seedlings were exposed to mild and severe drought stress, chlorophyll content was not affected (Ditmarová et al., 2009). Similar works with *Eucalyptus* spp. showed a chlorophyll content increase upon water stress (Correia et al., 2014b; Michelozzi et al., 1995). Carotenoids are involved in stress response as photoprotectors and ROS scavengers (Cazzonelli, 2011; Ismail et al., 2011). These natural pigments are also engaged in osmotic adjustment, preserving photosynthesis (Anjum et al., 2011). Furthermore they assist as precursors for abscisic acid biosynthesis (Cunningham Jr and Gantt, 1998).

Downregulation of photosystem II (PSII) activity gives rise to a discrepancy between the generation and use of electrons (Reddy et al., 2004). This effect can be analyzed by monitoring *in vivo* chlorophyll fluorescence. A method that depends upon the chlorophyll *a* fluorescence measurement in a dark adapted plant and after saturating light pulses (Schreiber, 2004). Such is a non-invasive and valuable tool for plant stress detection and photosynthesis assessment (Krause and Weis, 1991). Furthermore, as reviewed by Berger et al. (2007), chlorophyll fluorescence has proven effective in detecting photosynthetic disturbances before visible leaf damage occurred. However, the electron transport chain and its related processes are both resistant to drought stress (Ditmarová et al., 2009) and PSII activity may not decrease in a drought-dependent way (Sperdouli and Moustakas, 2012). Susiluoto and Berninger (2007) found an increased fluorescence in *Eucalyptus microtheca* under drought conditions and Correia et al. (2014b) had similar results with *E. globulus*. Down-regulation of effective photosystem II quantum yield was also found in several plant-pathogen interactions (Bonfig et

al., 2006; Chou et al., 2000; Swarbrick et al., 2006). Heat dissipation in the PSII core and antenna generate reactive oxygen species, leading to drought-induced oxidative stress (Reddy et al., 2004).

Oxidative response

Upon water stress-driven photosynthesis reduction, the excess excitation energy in photosystem II leads to a deficiency of the photosynthetic process with a consequent reactive oxygen species (ROS) accumulation (Aranjuelo et al., 2011; Chaves et al., 2009; Wilhelm and Selmar, 2011). Reactive oxygen species can react with DNA, lipids, and proteins, resulting in mutations and lipid/protein oxidation that lead to cell death and progressive organism aging (Beckman and Ames, 1998). ROS presence can be partially extrapolated through malondialdehyde (MDA) accumulation, a product of membrane lipid peroxidation (Cakmak and Horst, 1991). Plant pathogen recognition prompts a biphasic ROS accumulation with a rapid, but weak, transient phase, followed by a massive and prolonged second phase known as “oxidative burst”. Such is a front-line defense mechanism essential for further defense reactions (Heller and Tudzynski, 2011; Lamb and Dixon, 1997). Furthermore, ROS function as both localized and distance signals, promoting phytohormone accumulation, particularly, salicylic acid, and pathogenesis-related gene transcripts (Bostock et al., 2014; Naidoo et al., 2014).

Phytohormones

Phytohormones, besides controlling plant growth and development, also regulate the plant response to biotic and abiotic pressures (Bostock et al., 2014; Peleg and Blumwald, 2011). Several phytohormones are involved in the amplification of the initial defense signal, taking part in the exceptionally tailored response to the invader (Glazebrook, 2005; Naidoo and Za, 2013; Naidoo et al., 2011). Each of these signaling cascades encompasses a pathogenesis-related (PR) genes stimulation, intended to limit pathogen development and induce plant protection (Naidoo et al., 2011). Defense- or pathogenesis related (PR) proteins are an important part of the plant’s immune system, providing rapid host protection with low energy cost (Lay and Anderson, 2005).

Absciscic acid (ABA) is considered an universal hormone that regulates numerous developmental processes and adaptive stress responses in plants (Jiang and Hartung, 2008). This phytohormone is hoarded in root tissues and, upon water stress, is released into xylem vessels and transported to the shoot where, along with water potential reduction, induces stomatal closure (Jiang and Hartung, 2008; Rejeb et al., 2014). Such leads to a reduced water

loss via transpiration and, eventually, to a restriction of cellular growth (Peleg and Blumwald, 2011). Correia et al. (2014) and Jesus et al., (2015) found that, upon water stress, *E. globulus* plants present significantly higher levels of ABA. Absciscic acid has also been found to be a complex modulator of plant defense responses against phytopathogens (Denancé et al., 2013; Rejeb et al., 2014). It can act as a positive or negative regulator of plant defense, depending on the plant-pathogen interaction; it can have a negative interaction with salicylic and jasmonic acids under the combination of biotic and abiotic stresses (Denancé et al., 2013; Rejeb et al., 2014). Mauch-Mani and Mauch (2005) found that ABA can be implicated in increasing resistance of plants towards pathogens via its positive effect on callose deposition, an inducible defense that can prevent pathogen invasion (Rejeb et al., 2014).

Indole-3-acetic acid (IAA) is the predominant plant auxin involved in plant growth and development being found to promote stomatal opening (Peleg and Blumwald, 2011; Teale et al., 2006). Indole-3-acetic acid also plays a role in plant-pathogen interaction. Many plant pathogens can produce auxins or influence host auxin biosynthesis to interfere with the host's developmental processes (Wang et al., 2007). Nevertheless, plants developed mechanisms to repress auxin signaling during infection, as a defense strategy, particularly through salicylic acid production that, in turn, downregulates IAA production and signaling (Wang et al., 2007).

Salicylic acid (SA) is a phenolic compound and an important phytohormone found to be involved in several physiological processes such as seedling development, stomatal aperture, senescence, respiration and cell growth (Janda and Ruelland, 2015). It was also found to increase in content in response to drought (Hayat et al., 2010). For example, *Phillyrea angustifolia* and barley, showed an increase in SA concentration in response to drought (Bandurska, 2005; Munné-Bosch and Peñuelas, 2003). Pathogen infection leads to SA accumulation in uninfected leaves, suggesting an involvement in systemic acquired resistance (SAR) (An and Mou, 2011; Gaffney et al., 1993). Such is a mechanism that grants plant protection against a broad spectrum of pathogens at sites distant from the pathogen entry point (Janda and Ruelland, 2015). This hormone was identified as a main signaling molecule required for the plant's local defense responses against biotrophic pathogen attack (Denancé et al., 2013; Halim et al., 2006). Pathogen infection incites a SA levels increase in cells surrounding infection sites (Enyedi et al., 1992) to give rise to the hypersensitive response (HR). Such, is a locally triggered mechanism that leads to the collapse and death of infected cells (Alvarez, 2000). This cell death may control pathogens that infect living cells, but can fail to restrict necrotrophic pathogens (Govrin and Levine, 2000).

While biotrophic pathogen defense depends on programmed host cell death, necrotrophic pathogens benefit from that mechanism and are, thus, targeted by the induction of the jasmonic acid pathway (Glazebrook, 2005). Jasmonic acid (JA), and its metabolites, collectively known as jasmonates, were found to be involved in plant responses to abiotic as well as to biotic stresses (Ghasemi Pirbalouti et al., 2014). Jasmonic acid was found to have an effect on abiotic stress in strawberry (Wang, 1999), pear (Gao et al., 2004) and *E. globulus* (Correia et al., 2014b). Furthermore, the JA/ET pathway is also involved in the stimulation of the induced systemic resistance (ISR) (Naidoo et al., 2014). Such results in priming, the induction of cellular defense responses in tissues distal to the point of stimulation (Pozo et al., 2008).

Induced responses to pathogens: A crosstalk mechanism

The first line of defense against phytopathogens is preformed (constitutive). It is present in host cells prior to pathogen contact, being considered a nonspecific shield (Oghenekaro et al., 2013). Once preformed defenses are breached, induced responses are activated (Jones and Dangl, 2006). These require recognition of non-self-cells by general elicitors called PAMPs (Pathogen Associated Molecular Patterns) or MAMPs (Microbe Associated Molecular Patterns), subsequent signaling and production of defensive products (Naidoo et al., 2014; Trouvelot et al., 2014). These patterns (MAMPs and PAMPs) are detected by pattern recognition receptors (PRRs), that lead to the non-specific immune response known as “pattern-triggered immunity” (PTI) (Dardick and Ronald, 2006; Morkunas and Ratajczak, 2014; Naidoo et al., 2014). Such immunity mechanism depends on MAP kinase activation, ROS production, transcriptional reprogramming, hormone biosynthesis and deposition of callose in the cell wall (Jones and Dangl, 2006; Morkunas and Ratajczak, 2014). According to some studies, cross-species PRRs could be potentially used to improve *Eucalyptus* spp resistance, being, therefore, attractive targets for manipulation (Naidoo et al., 2014).

Some pathogens have developed strategies to avoid or even suppress the plant’s PTI (Denancé et al., 2013). To counteract this, a second line of defense, more specific, rapid, and effective, known as “effector-triggered immunity” (ETI) takes action (Dodds and Rathjen, 2010). It depends of the recognition of effectors by plant resistance proteins (PRP) (Bolouri and Van den Ende, 2012; Naidoo et al., 2014) and involves mechanisms such as oxidative burst and hypersensitive response, being associated with extensive changes in gene expression (Naidoo et al., 2014). With the recent knowledge development on both processes, it has become more and more clear that both PTI and ETI are probably regarded as two

mechanisms of the same process (Boller and Felix, 2009). Nonetheless, this subject is still in its beginnings and additional studies should be performed.

The master dissertation: main purposes

The current climate change scenario, with its high extent and speed, is projected to induce direct growth impairment as well as hazardous changes in plant-pathogen interactions. Such is expected to increase disease incidence and severity, particularly to *Eucalyptus* spp. plantations. Considering the fundamental importance of *E. globulus* and how drought increases the threat that *N. kwambonambiense* poses to plantations, it is of major concern to assess the predisposition effect that drought may have in *N. kwambonambiense* infection in order to provide insights that can be of service to the eucalypt forest industry.

The aim of this study was, on a secondary level, to evaluate the *Eucalyptus globulus* response to the individual drought and pathogen stresses and, on a primary level, to assess the predisposition effect that drought stress may have on *Eucalyptus globulus* infection by *Neofusicoccum kwambonambiense*. To achieve these objectives, a holistic set of morphological, physiological, and biochemical traits were assessed, including fungal progression analysis, water relations, osmolyte accumulation, lipid peroxidation and photosynthetic and hormonal responses.

The master thesis disclosed herein is presented as a research paper and it is prepared according to the authors' instructions from the journal Environmental and Experimental Botany.

Part II

Physiological and hormonal changes behind *Eucalyptus* predisposition to *Neofusicoccum kwambonambiense* under water stress

Glória Pinto^{1*}, Helder Berenguer^{1*}, Barbara Correia¹, Aurelio Gómez-Cadenas², Andreia Cerqueira¹, Artur Alves^{1#}

¹ Department of Biology & CESAM – Centre for Environmental and Marine Studies, University of Aveiro, Campus Universitário de Santiago, 3810-193 Aveiro, Portugal

² Universitat Jaume I, Departamento de Ciencias Agrarias y del Medio Natural, 12071 Castelló de la Plana, Spain

*Authors with equal contribution

corresponding author: artur.alves@ua.pt; tel +351 234 247 182; fax + 351 234 372 587

Abstract

Eucalyptus spp. plantation forestry has exponentially grown and currently represents a valuable economic commodity through the paper pulp and timber industry. *Neofusicoccum kwambonambiense* is an endophytic opportunist plant pathogen known to severely affect *Eucalyptus globulus*. Global temperatures have been increasing and this trend is expected to maintain in the future with expected losses to the eucalypt industry, directly, and as the result of the interaction between pathogens and climate change-driven drought. Taking such in consideration, it is of major importance to assess, on a physiological and hormonal levels, the predisposition effect that drought may have in *N. kwambonambiense* infection, in order to provide insights that may be of service to the eucalypt forest industry. To do so, *E. globulus* plants were inoculated with *N. kwambonambiense* after a 23-day water stress period (18% of field capacity). Physiological and biochemical parameters and hormonal content were assessed 65-days post-inoculation. Water-stress lead to a water potential decrease and a further ABA-triggered stomatal closure. Although hormonal (salicylic and jasmonic acid), oxidative and high-sugar-resistance responses took place, the plant was unable to properly react to oxidative damages (observable through malondialdehyde content) that, along with the already imposed gas restrictions, lead to a decrease in the photosynthetic efficiency. Such resulted in a compromising cycle of increased lesion extension and further susceptibility to infection. Ultimately a higher *N. kwambonambiense* progression and disease expression arose, therefore substantiating the predisposition hypothesis. This work fills a scientific gap, not only verifying the water stress predisposition effect but also exploring the plant's physiological response, unveiling plant fitness and its role on water stress predisposition.

Keywords: Forest trees, Plant performance, Biotic and abiotic interactions, Plant-pathogen interactions

Introduction

Forest tree growth, productivity and survival can be negatively affected by environmental stresses (Osakabe et al., 2012). Their effects are expected to become more pronounced in both duration and severity in the near future, partly do climate change and the occurrence of simultaneous stress factors (FAO, 2015; Stella et al., 2013), particularly across Europe and the Mediterranean region (Blenkinsop and Fowler, 2007). Furthermore, according to several studies, water availability has been shown to be the main factor limiting plant growth, particularly of eucalypt plantations (Shvaleva et al., 2006; Villar et al., 2011).

Eucalyptus spp., known for their rapid growth, reduced rotation time and high adaptability to soils and climate, have become a key genera amongst planted hardwoods (Booth, 2013; Florence, 2004; Luger, 2003). Native to Australasia, they are explored in more than 90 countries, with main plantations located in Brazil, India and China, totaling more than 20 million ha (Brooker and Hopper, 2002; Iglesias-Trabado and Wilstermann, 2008). They represent a key source of biomass for fibreboard, industrial charcoal, fuelwood and paper pulp, being likewise recognized as a potential biofuels and biomaterials source (Hinchee et al., 2011; Luger, 2003; Turnbull, 1999). According to the most recent Portuguese National Forest Inventory (ICNF, 2013), *Eucalyptus* spp., particularly *Eucalyptus globulus* (Labill), occupies more than 800 000 planted ha, corresponding to 26% of the total forest area. *E. globulus* was introduced in Portugal around 1860 for timber and paper pulp production (Borrallho et al., 2007) and, due to their high growth rates, pulp yield and environmental adaptability (Costa et al., 2004), have become the dominant forest type with a considerable economic importance through the paper pulp industry (CELPA, 2014).

Neofusicoccum kwambonambiense (Pavlic, Slippers & M.J) is part of the *Botryosphaeriaceae* spp. (*Botryosphaeriales*, Ascomycetes), a vast distribution family that occurs on a wide range of dicotyledonous, monocotyledonous and gymnosperm hosts (Crous et al., 2006; Slippers and Wingfield, 2007). *Botryosphaeriaceae* spp. are mostly considered opportunistic endophytes that, upon stressful conditions, such as hail, frost, insect damage, and, particularly, drought, adopt a pathogenic behavior (Crous et al., 2006). When affecting *Eucalyptus* spp., lead to shoot and branches dieback, stem cankers followed by kino exudation, coppice failure, and, in severe cases, tree death (Slippers et al., 2009; Slippers and Wingfield, 2007). Their high occurrence, dispersion aptitude and ability to cause disease when their hosts are under stress, makes this pathogen a significant threat to native forests and plantations, specially *Eucalyptus* spp., where they are explored as non-native species (Crous et al., 2006; Slippers et al., 2004; Slippers and Wingfield, 2007). Barradas et al. 2016 detected,

in Portugal, in 2012, the first recorded occurrence of *N. kwambonambiense* in *E. globulus*, which was considered the most aggressive species amongst the other *Botryosphaeriaceae* species found.

There is a growing acknowledgement amongst the scientific community and policy makers that sustainable forest management is affected by multiple factors associated with global climate change (Noss, 2001). Although under field conditions, often two or more stresses are simultaneously or successively associated (Chelli-Chaabouni, 2014) most of the research has largely focused on individual biotic or abiotic stresses (Niinemets, 2010). Several studies regarding the effect of water stress on *Eucalyptus* spp. have been explored (Correia et al., 2016, 2014b; Li et al., 2000; Martorell et al., 2014; White et al., 1996a). Additionally, the distribution and effect of *Botryosphaeriaceae* on *Eucalyptus* spp. has also been addressed (Alves et al., 2013; Barradas et al., 2016; Pavlic et al., 2007; Pérez et al., 2010; Slippers and Wingfield, 2007). Nonetheless, the response to the collective drought and pathogen stress, from a predisposition point of view, cannot be predicted based in the plant's response to the individual stresses, being assured to lead to a more complex picture, particularly, under a climate change scenario (Desprez-Loustau et al., 2006; Niinemets, 2010; Rejeb et al., 2014).

The role of water stress has been object of predisposition-related research (Bostock et al., 2014; Desprez-Loustau et al., 2006). The predisposition phenomenon is the consequence of abiotic stresses occurring prior to infection, increasing plant disease susceptibility (Bostock et al. 2014). Furthermore, when hosts are subjected to drought preceding inoculation, symptom development tends to be more severe (Jactel et al., 2012; Van Niekerk et al., 2011). Forest-related studies regarding predisposition dealt with different pathosystems, including: *Quercus* spp.- *Botryosphaeria stevensii* and *Phytophthora cinnamomi* (Luque et al., 2002), *Populus nigra* × *P. maximowiczii* - *Septoria musiva* (Franceschini et al., 2004), *Pinus resinosa* - *Sphaeropsis sapinea* (Blodgett and Stanosz, 1997), *Eucalyptus* spp. - *Botryosphaeria dothidea* (Smith et al., 1994) and grapevines - *Botryosphaeriaceae* spp. (Van Niekerk et al., 2011), and conclusively reported increased susceptibility and symptom severity under predisposing water stress conditions.

The integration of plant responses to abiotic and biotic stress requires the involvement of multiple strategies and signals that may be common or specific to each type of stress (Bostock et al., 2014). The degree of plant susceptibility to pathogen attack depends on the host condition upon infection but also on the host's metabolic status, crucial for pathogen growth, as it appears that pathogens are capable to manipulate different aspects of plant

metabolism to achieve optimal growth conditions (Chen et al., 2010; Desprez-Loustau et al., 2006; Dodds and Rathjen, 2010). Adaptation to both abiotic and later biotic stress have an impact on primary metabolism homeostasis (Bolton, 2009; Chaves et al., 2003). In fact, defense-related components synthesis (metabolites and proteins) is energy demanding (Bolton, 2009), and may be compromised since drought potentially leads to a disturbance in the energy balance as stomatal conductance takes place and photosynthesis is reduced (Correia et al., 2016, 2014b; Desprez-Loustau et al., 2006). Stomatal closure depends on the accumulation of solutes and the phytohormone abscisic acid (ABA) (Correia et al., 2014b; Jesus et al., 2015). This point represents a node of convergence in the crosstalk between water stress and changes in disease susceptibility as ABA, a primary dehydration-responsive messenger (Jiang and Hartung, 2008), has been found to increase susceptibility or resistance to different pathogens (Mauch-Mani and Mauch, 2005; Rejeb et al., 2014). Such dual effect makes ABA a divisive molecule, whose effects depends of the environmental conditions (Luna et al., 2011). Furthermore, ABA may operate, or not, through an antagonistic relationship with important disease resistance signals such as salicylic acid, and jasmonic acid (Bari and Jones, 2009; Peleg and Blumwald, 2011). Reactive oxygen species (ROS) signaling (Miller et al., 2008) together with plant hormones therefore represent major components of the plant's response regulatory networks, underlying abiotic stress adaptation and pathogen recognition (Bostock et al., 2014; Peleg and Blumwald, 2011).

Increased losses to the eucalypt industry are expected, as a result of the interaction between climate change-driven drought and pathogen increased incidence (Sturrock et al., 2011). Taking into consideration *E. globulus* economic benefits, climate change effects on forest pathogens, particularly on *N. kwambonambiense* dynamics, this study intends to explore how water limitation affects *E. globulus* susceptibility to *N. kwambonambiense*. Physiological and hormonal changes will be assessed to unveil plant fitness and its role on water stress predisposed plants. By predicting a situation that under a climate change scenario may worsen a problem already encountered by forests, we aim to contribute to a better forest management and sustainability.

Materials and methods

Plant, fungal material and growing conditions

Neofusicoccum kwambonambiense (strain CAA755) (Fig. 1) belonging to A. Alves culture collection, Universidade de Aveiro, Portugal, was isolated from *E. globulus* stem cankers, maintained on ½ strength potato-dextrose-agar (PDA) medium (Merck, Germany) at environmental temperature (about 20–25 °C). Strains were subcultured one-week prior to inoculation.

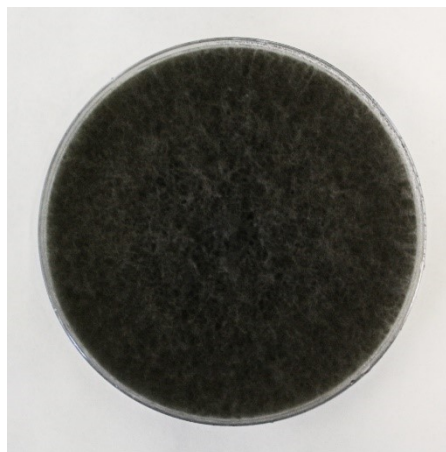


Figure 1 *N. kwambonambiense* grown on ½ strength PDA for one week

Four month old, *E. globulus* (clone GM-2-58) rooted cutting were obtained from Altri Florestal SA (Portugal), transplanted to 2 L plastic pots and filled with 3:2 (w/w) peat:perlite mixture. Potted cuttings were subjected to a 66-day acclimation period, watered on a regular basis to maintain 80% of field capacity (FC) and fertilized once a week with a (N:P:K 12:4:6) nutritive solution (FRUTIFOL L12®). To minimize the effects of environmental heterogeneity, pots were randomly arranged and sporadically repositioned to neighboring positions during the experiment. Pot weight was monitored gravimetrically. During the experimental period, daily mean air temperatures showed a progressive increase from 5-14°C (min.-max. temp.) in February, 2015 to 15-23°C (min.-max. temp.) at the end of the experiment, in July, 2015.

Experimental design

Plants were assigned to different water regime and fungi presence treatments (Fig. 2). Ten plants each made up the non-inoculated group, either well-watered (WW-NF), or water-stressed (WS-NF). Eighteen plants each made up the fungi-inoculated group, either well-watered (WW-F), or water-stressed (WS-F). WW plants were maintained in the same conditions of the acclimation period and WS plants were assigned to a water deficit regime of 18% FC. Such was achieved at the end of a 23-day period, by gradually reducing water supply. Once water-stressed was achieved, WW-F and WS-F plants were inoculated with *N. kwambonambiense*. To do so, a superficial wound, 1.5 cm long, was made on the stem of every plant with a sterile blade, 3-5 cm above ground level. Colonized PDA plugs, fungus-side-down, were placed on the wounds and immediately sealed with Parafilm® (American National Can, Greenwich, CT, USA). WW-NF and WS-NF plants stems received non-colonized PDA plugs. Plants were kept in these conditions throughout 65 days. To determine the time to end

the experiment, periodical external lesion extension measurements were made to identify the moment at which WS-F plants presented significantly higher values than WW-F plants.

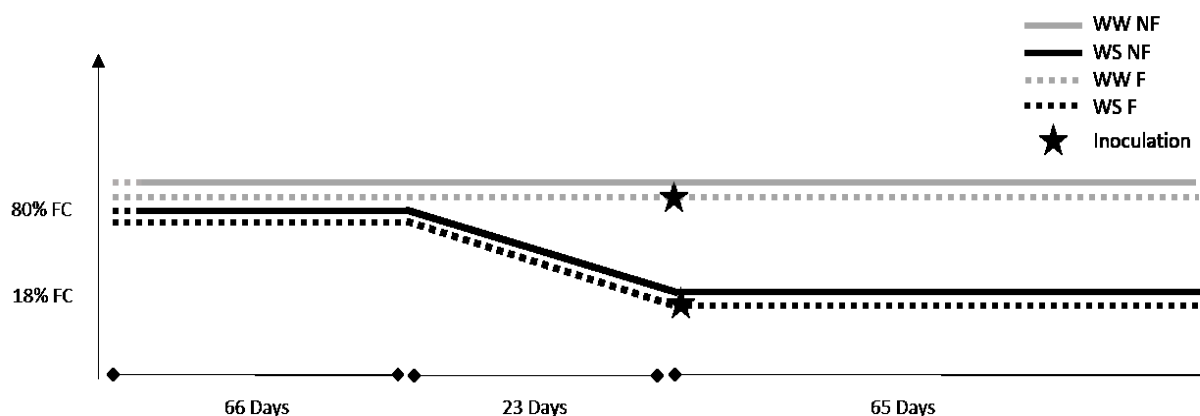


Figure 2 Experimental design implemented to evaluate the response of WW-NF, WS-NF, WW-F and WS-F, *E. globulus*. WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

Sample collection

Upon inoculation and till the end of the experiment, height and external lesion extension were periodically measured. At the end of the experiment (65 days-post inoculation (DPI)), the following *in situ* parameters were recorded in plants of each treatment group: external lesion extension, height, water potential, leaf gas-exchange and stomatal conductance parameters and chlorophyll *a* fluorescence. Also, fully expanded leaves were harvested and immediately frozen into liquid nitrogen for further biochemical analysis: total soluble sugars (TSS) and proline content, estimation of lipid peroxidation through malondialdehyde (MDA) quantification, chlorophylls *a,b* and carotenoid content and hormone quantification.

Height, external lesion extension and confirmation of *N. kwambonambiense* presence

Height (in cm), measured from the substrate surface to the apical meristem, was assessed in all plants. External lesion extension (in cm) was measured in all fungi-inoculated plants (both WW-F and WS-F). To confirm *N. kwambonambiense* presence, and confirm Koch's postulates, stem portions were extracted from the region where the lesion was made (L), in the intermediate point between the lesion and plant apex (L+) and near the plant apex (L++). Isolations were made by directly plating out stem portions on PDA medium (Merck, Germany) after surface sterilization. Plant tissue was sterilized by placing it in 5% sodium hypochlorite for 1 min, followed by 96 % ethanol for 1 min and rinsed in sterile water for 1 min. Plates were incubated at room temperature and checked regularly for fungal presence and

growth. Non-inoculated plant wounds (both WW-NF and WS-NF) successfully healed and no fungus was isolated from them.

Water status

Midday shoot water potential (Ψ_{md}) was measured in six randomly selected plants per treatment group with a Scholander-type pressure chamber (PMS Instrument Co., Corvallis, OR).

Leaf gas-exchange and stomatal conductance parameters

Intercellular CO₂ concentration content (C_i , ppm), stomatal conductance (g_s , mol H₂O m⁻² s⁻¹), transpiration rate (E , mmol H₂O m⁻² s⁻¹), and net CO₂ assimilation rate (A , μ mol CO₂ m⁻² s⁻¹) were performed using a portable infrared gas analyser (LCpro-SD, ADC BioScientific Ltd., UK) equipped with a broad leaf chamber. To determine the saturation light intensity A/PPFD (photosynthetic photon flux density; light response curves of CO₂ assimilation) curves were performed with the following PPFD: 2000, 1500, 1000, 750, 500, 250, 100, 50 and 0 μ mol m⁻² s⁻¹. After A/PPFD data analysis, punctual measurements at saturation light intensity were performed at 1500 μ mol m⁻² s⁻¹. The following conditions were maintained inside the chamber during all the measurements: air flux: 200 mol s⁻¹; block temperature: 25 °C; and atmospheric CO₂ and H₂O concentration. Data were recorded when the measured parameters were stable (2–6 min).

Chlorophyll a fluorescence and photosynthetic pigments quantification

Steady-state modulated chlorophyll fluorescence was determined with a portable fluorometer (Mini-PAM; Walz, Effeltrich, Germany). Light adapted components of chlorophyll fluorescence were measured: steady-state fluorescence (F), maximal fluorescence (F'_m), variable fluorescence F'_v (equivalent to $F'_m - F$) and quantum yield of PSII photochemistry (ϕ PSII) equivalent to $(F'_m - F)/F'_m$. Leaves were then dark adapted for no less than 20 min to obtain F_0 (minimum fluorescence), F_m (maximum fluorescence), F_v (variable fluorescence, equivalent to $F_m - F_0$) and F_v/F_m (maximum quantum yield of PSII photochemistry).

Chlorophylls *a* and *b* and carotenoids content were quantified according to Sims and Gamon (2002). Pigments were extracted with acetone/Tris (50 mM) buffer at pH 7.8 (80/20) (v/v). After homogenization and centrifugation, supernatants were used to read absorbances at 663 nm, 537 nm, 647 nm and 470 nm (Thermo Fisher Scientific Spectrophotometer, Genesys 10-uv S, Waltham, MA). Chlorophyll *a* was calculated as $(0.01373 A_{663} - 0.000897 A_{537} - 0.003046 A_{647})$ and chlorophyll *b* as $(0.02405 A_{647} - 0.004305 A_{537} - 0.005507 A_{663})$.

Carotenoids content was calculated as $[(A_{470} - (17.1 \times (Chl_a + Chl_b) - 9.479 \times B)) / 119.26]$, where $B = (0.08173 A_{537} - 0.00697 A_{647} - 0.002228 A_{663})$.

Lipid peroxidation

Lipid peroxidation on leaves was estimated by measuring the amount of malondialdehyde (MDA) which takes into consideration the possible effects of interfering compounds in the assay for thiobarbituric acid (TBA)-reactive substances using the method described by Hodges et al. (1999) with slight changes. Briefly, MDA was extracted from 50 mg frozen leaves using 1.25 mL of 80% (v/v) ethanol. After centrifugation, the supernatant was mixed with either 1 mL of 20 % TCA + 0.01% BHT and 20 % TCA + 0.5% TBA + 0.01% BHT and heated at 98 °C for 30 m. Afterwards, samples were placed on ice for 5 m and absorbance was read at 600 nm, 532 nm and 440 nm against control. MDA equivalents (nmol mL⁻¹) were calculated as $(A-B/157\ 000) \times 10^6$, where $A = [(Abs_{532} + TBA) - (Abs_{600} + TBA) - (Abs_{532} - TBA - Abs_{600} - TBA)]$, and $B = [(Abs_{440} + TBA - Abs_{600} + TBA) \times 0.0571]$.

Total soluble sugars and proline quantification

Total soluble sugars (TSS) content was determined by the anthrone method, as described by Irigoyen et al (1992). Briefly, TSS were extracted from 100 mg frozen leaves using 10 mL of 80 % (v/v) ethanol at 80 °C for 1 h. After centrifugation, supernatant was mixed with 1.5 mL of anthrone and incubated at 100 °C throughout 10 min. Absorbance was read at 625 nm and TSS content was calculated against a D-glucose standard curve (0 to 1.25 mg/ml).

Proline content was determined as described by Khedr et al. (2003), with slight modifications. Briefly, about 100 mg of frozen plant material were homogenized in 1.5 ml of 3% sulphosalicylic acid, shaken vigorously and centrifuged at 10,000 g for 10 min at 4 °C. 1 ml of extract was incubated with 1 ml of ninhydrin acid and 1 ml of glacial acetic acid for 1 h at 100 °C. Afterwards, the reaction mixture was cooled in ice for 20 min. 1 ml of toluene was added and the mixture and left for 30 min until phases separation. Absorbance of the toluene phase was measured at 520 nm. Proline concentration was determined against a L-proline standard curve (0 to 50 µmol mL⁻¹).

Hormone quantification

Hormone extraction and analysis were carried out following the procedure described in Durgbanshi et al. (2005) with slight modifications. Frozen fresh tissue (0.2 g) was spiked with 100 ng of d6-ABA, 100 ng of Dihydrojasmonic acid, 5 ng of [2H₂] IAA, and 100 ng of d6-SA, prepared as in Gomez-Cadenas et al. (2002) and homogenized with 5 ml of distilled water.

After centrifugation at 4000 x g at 4 °C, supernatants were recovered and pH adjusted to 3 with 30 % acetic acid. The acidified water extract was partitioned twice against 3 ml of diethyl ether. The organic upper layer was recovered and vacuum evaporated in a centrifuge concentrator (Speed Vac, Jouan, Saint Herblain Cedex, France). The dry residue was then resuspended in a 10 % MeOH solution by gentle sonication. The resulting solution was passed through 0.22 µm regenerated cellulose membrane syringe filters (Albet S.A., Barcelona, Spain) and directly injected into a UPLC system (Acquity SDS, Waters Corp., Milford, MA, USA).

Analytes were separated by reversed-phase (Nucleodur C18, 1.8 µm 50 × 2.0 mm, Macherey- Nagel, Barcelona, España) using a linear gradient of ultrapure H₂O (A) and MeOH (B) (both supplemented with 0.01 % acetic acid) at a flow rate of 300 µl min⁻¹. The gradient used was: (0-2 min) 90:10 (A:B), (2-6 min) 10:90 (A:B) and (2-6-7 min) 90:10 (A:B). Hormones were quantified with a Quattro LC triple quadrupole mass spectrometer (Micromass, Manchester, UK) connected online to the output of the column through an orthogonal Z-spray electrospray ion source. Quantitation of plant hormones was achieved by external calibration with standards of known amount.

Statistical analysis

Student's t-test (t-test) was used to evaluate the effect of watering regime on the lesion extension of fungi inoculated plants (WW-F and WS-F). Two-way analysis of variance (ANOVA) was used to assess the effects of watering regime, fungi treatment and their interaction, on height, water potential, TSS, proline, gas exchange (c_i , g_s , E, A), MDA, pigments (Chl *a*, *b* and carotenoids), chlorophyll *a* fluorescence and hormone content (ABA, IAA, JA, SA). Whenever an interaction between watering regime and fungi treatment was found, a simple main effect analysis (ANOVA one-way) was carried (using the error term of the two-way ANOVA as proposed by Quinn and Keough, (2002)). When applicable, Holm-Sidak multiple comparison tests were employed to identify significant differences between treatments. T-test's, two-way ANOVAS and Holm-Sidak tests were performed using SigmaPlot (SigmaPlot for Windows v. 12.0, Systat Software Inc.) and main effects were calculated with MS Excel (Microsoft, Redmond, WA). Prior to analysis, data were ln- or 1/x-transformed to correct heteroscedasticity. (*) represent significant differences within water regimes, and (#) represent significant differences within fungi treatments All analyses used a significance level of 0.05.

Principal components analysis (PCA) was carried out to explore the morpho-physiological profile of *E. globulus* plants subjected to the different treatments by reducing the

multivariate data matrix to an interpretable bidimensional biplot that explains the highest proportion of variation in the data (ter Braak and Verdonschot, 1995). External lesion extension was excluded from the PCA because this parameter wasn't evaluated on non-fungi inoculated plants. PCA was conducted with Minitab 17 Statistical Software (2010). State College, PA: Minitab, Inc. Available from www.minitab.com.

Results

Height, external lesion extension and *N. kwambonambiense* presence confirmation

Water regime was the main influence on height differences (Fig. 3) (see F-ratios in Table 3) in a way that F plants did not differ from NF plants while WW plants consistently showed superior heights when compared with WS plants. At a morphological level, WS-F plants, besides presenting thinner stems, also presented larger internal lesion extensions (Fig. 4a). On an external level (Fig. 4b), at 58 DPI, WS-F external lesion extension was significantly larger than WW-F. To confirm whereas such difference was sustained, plants were kept for another 7 days, at which time, significant differences were maintained and the experiment came to an end. Moreover, *N. kwambonambiense* consistently reached superior heights in WS-F plants, 75% vs. 25%, for the L+ region and 25% vs. 0% for the L++ region (Table 1).

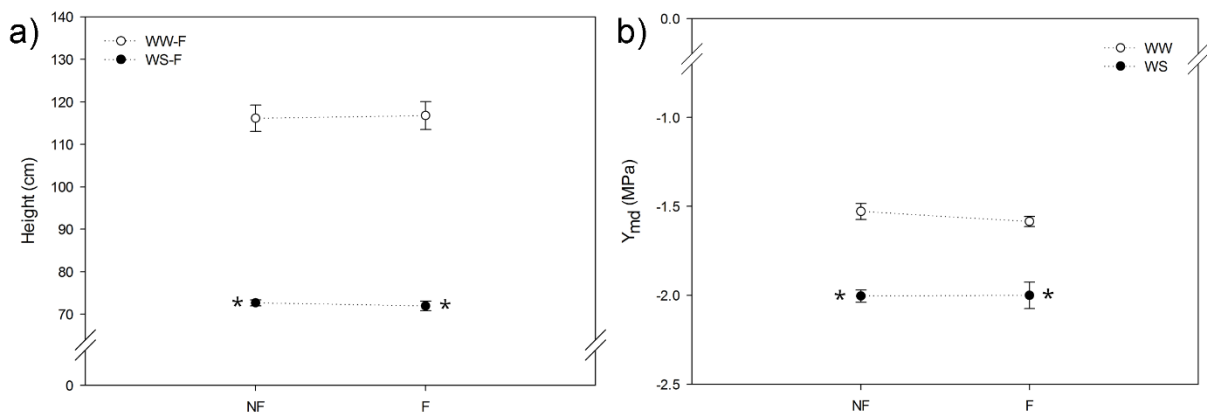


Figure 3 Height (a) and midday water potential (b) of WW-NF, WS-NF, WW-F and WS-F *E. globulus*. Data are presented as mean \pm SE. (*) indicate significant differences within water regimes and (#) indicate significant differences within fungi treatments. ($p \leq 0.05$). WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

Table 1 Fungi presence in WW-F and WS-F *E. globulus* at different plant regions. In the region where the lesion was made and the fungi inoculated (L), between the lesion and plant apex (L+) and near the plant apex (L++). The presence of the fungi is shown in percentage of a total of 8 plants per treatment. WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

Region/ Treatment	WW-F	WS-F
L++	0%	25%
L+	25%	75%
L	100%	100%

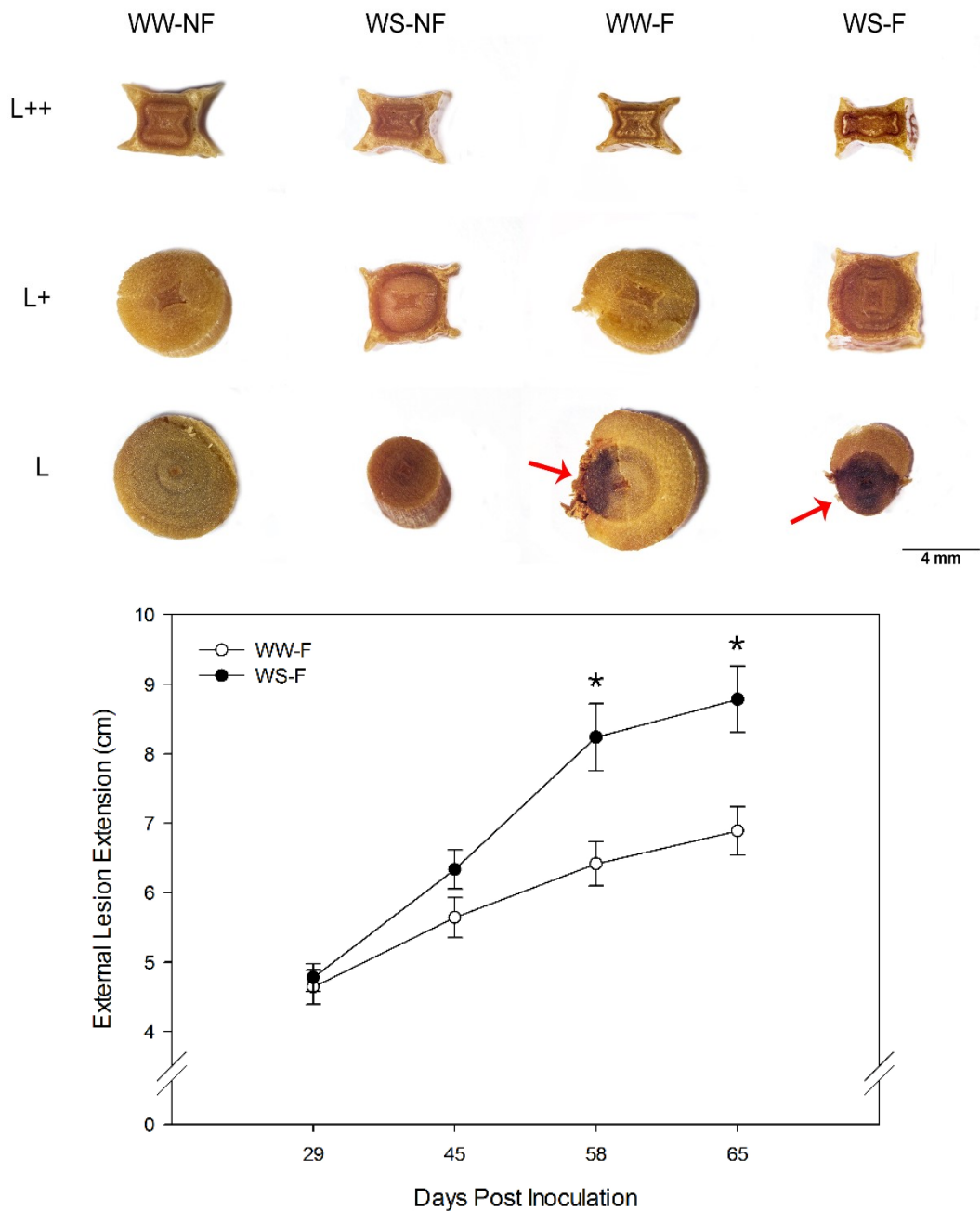


Figure 4 (a) Morphological aspect of WW-NF, WS-NF, WW-F and WS-F *E. globulus* stem cuts at different regions. In the region where the lesion was made and the fungi inoculated (L), between the lesion and plant apex (L+) and near the plant apex (L++). Red arrows indicate lesion affected areas. (b) External lesion extension of WW-F and WS-F *E. globulus*. Data are presented as mean \pm SE. (*) indicate significant differences within water regimes and (#) indicate significant differences within fungi treatments. ($p \leq 0.05$). WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

Water status

Water regime was the main influence on water potential differences (Fig. 3b) (see F-ratios in Table 3) in a way that F plants did not differ from NF plants while WW plants consistently showed superior water potentials when compared with WS plants.

Leaf gas-exchange and stomatal conductance parameters

A significant interaction between water regime and fungi treatment was found in all gas exchange parameters (Fig. 5). C_i (Fig. 5a) presented a decrease upon water stress in NF plants and an increase in WS plants caused by *N. kwambonambiense* presence. Regardless of fungi presence, water stress caused a decrease in g_s (Fig. 5b), E (Fig. 5c) and A (Fig. 5d) while an increase can be observed in WW plants as a result of interaction with *N. kwambonambiense*. Presence of *N. kwambonambiense* caused a decrease of E (Fig. 5c) and A (Fig. 5d) in already water-stressed plants. In all parameters, water regime was the main influence (see F-ratios in Table 3).

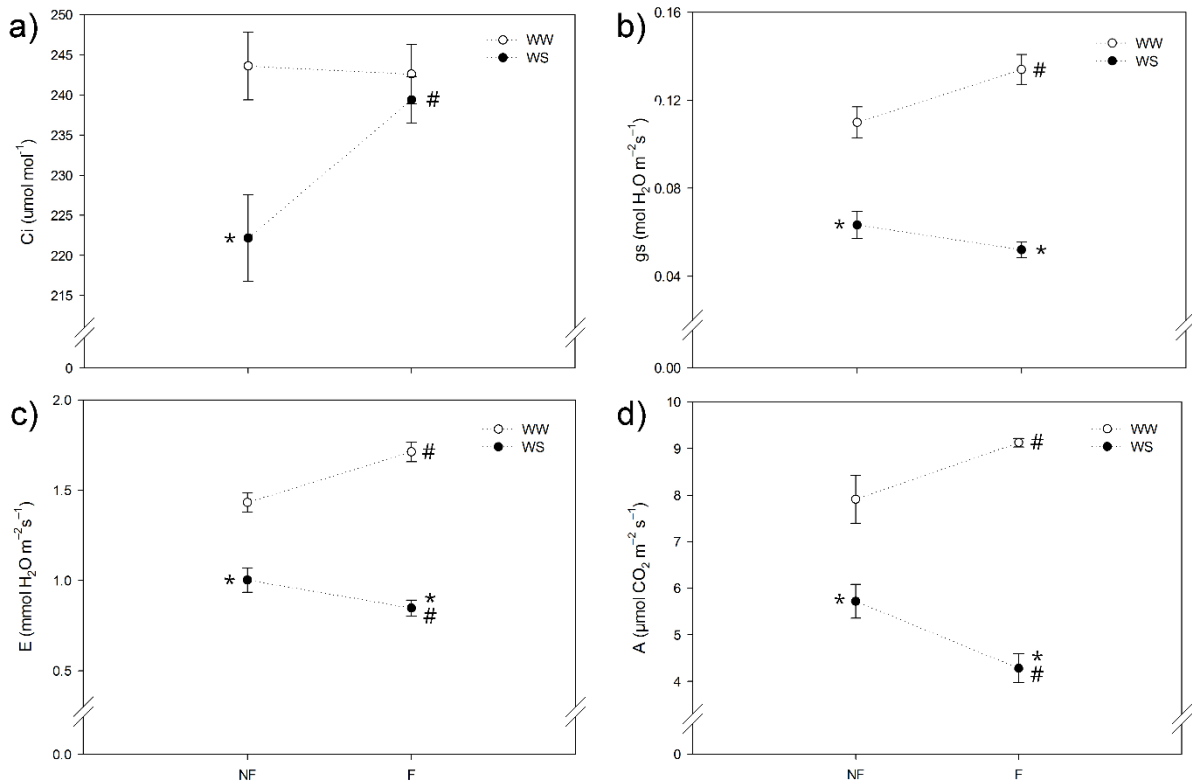


Figure 5 Intercellular CO_2 concentration (c_i) (a), stomatal conductance (g_s) (b), transpiration rate (E) (c), and foliar net CO_2 assimilation rate (A) (d) of WW-NF, WS-NF, WW-F and WS-F *E. globulus*. Data are presented as mean \pm SE. (*) indicate significant differences within water regimes and (#) indicate significant differences within fungi treatments. ($p \leq 0.05$). WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

Chlorophyll a fluorescence and photosynthetic pigments quantification

No significant differences regarding Ch *a*, *b* and carotenoids concentration or F_v/F_m were found (Table 2). Only $\Phi PSII$ (Table 2) presented differences, with an increase in WW plants caused by *N. kwambonambiense* and a decrease in F plants caused by water stress. In this case, fungi presence was the main influence (see F-ratios in Table 3).

Table 2 Chlorophyll *a,b* and carotenoids content, maximum quantum yield (F_v/F_m) and effective quantum yield ($\Phi PSII$) of WW-NF, WS-NF, WW-F and WS-F *E. globulus*. Data are presented as mean \pm SE. (*) indicate significant differences within water regimes and (#) indicate significant differences within fungi treatments. ($p \leq 0.05$). WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

Treatment	Chl <i>a</i> content ($\mu\text{mol g}^{-1}\text{DW}$)	Chl <i>b</i> content ($\mu\text{mol g}^{-1}\text{DW}$)	Carotenoids content ($\mu\text{mol g}^{-1}\text{DW}$)	F_v/F_m	$\Phi PSII$
WW-NF	1.570 ± 0.124	0.576 ± 0.037	0.728 ± 0.045	0.823 ± 0.003	$0.742 \pm 0.002 \#$
WS-NF	1.574 ± 0.077	0.578 ± 0.079	0.691 ± 0.037	0.824 ± 0.003	0.748 ± 0.002
WW-F	1.760 ± 0.116	0.640 ± 0.051	0.815 ± 0.064	0.823 ± 0.001	$0.760 \pm 0.004\#^*$
WS-F	1.738 ± 0.076	0.640 ± 0.027	0.742 ± 0.027	0.826 ± 0.002	$0.749 \pm 0.004 \#^*$

Lipid peroxidation

A significant interaction between water regime and fungi treatment was found MDA content (Fig. 6a). Under WW conditions, *N. kwambonambiense* led to a decrease in MDA content. The effect was opposite in WS conditions, with an increase. A similar increase was identified in F plants under water stress conditions. In this case, water regime was the main influence (see F-ratios in Table 3).

Total soluble sugars and proline quantification

A significant interaction between water regime and fungi treatment was found in TSS (Fig. 6b) and proline content (Fig. 6c). Presence of *N. kwambonambiense* led to an increase of TSS in WW and WS conditions, although heightened in WS conditions. Accordingly, fungi treatment was the main influence (see F-ratios in Table 3). Presence of *N. kwambonambiense* led to an increase of proline content in WW conditions, as did water stress in NF plants. Water regime was, again, the main influence (see F-ratios in Table 3).

Hormones

Regardless of *N. kwambonambiense* presence, water stress caused an increase in ABA content (Fig. 7a). In IAA content (Fig. 7d), a decrease is observed in NF plants. In both ABA and IAA content, water regime was the main influence (see F-ratios in Table 3). Water stress caused a decrease of JA (Fig. 7b) content in NF plants while in WS conditions, fungi inoculation led to an increase. Furthermore, a significant interaction between water regime and fungi treatment can be seen in JA content. In all treatments, both water stress and *N. kwambonambiense* caused an increase in SA

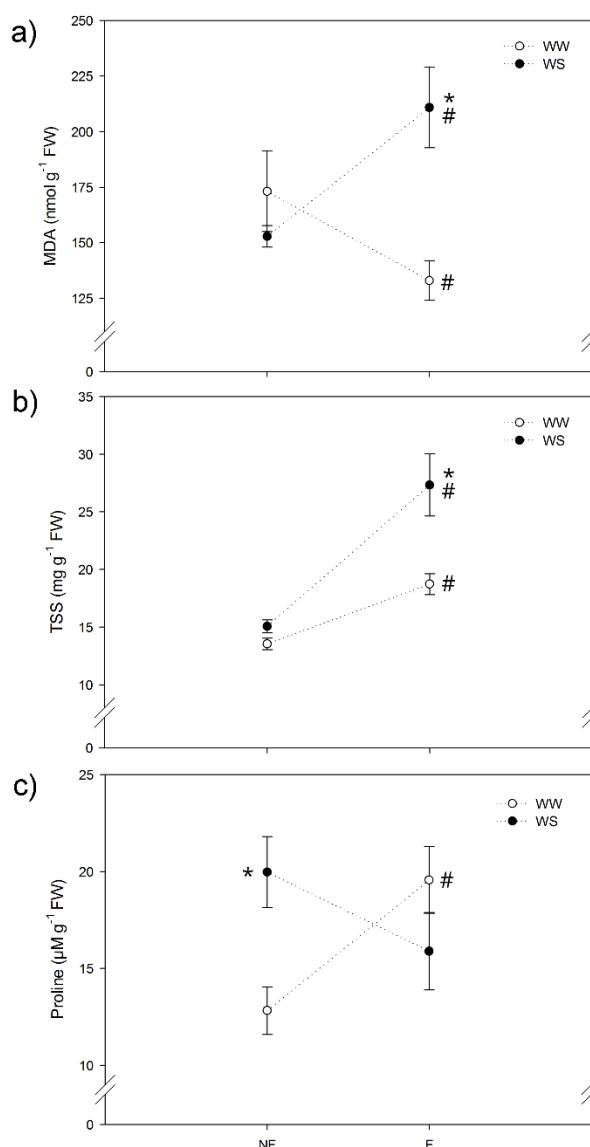


Figure 6 MDA (a), TSS (b) and proline (c) content in WW-NF, WS-NF, WW-F and WS-F *E. globulus*. Data are presented as mean \pm SE. (*) indicate significant differences within water regimes and (#) indicate significant differences within fungi treatments. ($p \leq 0.05$). WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

content (Fig. 7c). In both JA and SA content, fungi presence was the main influence (see F-ratios in Table 3).

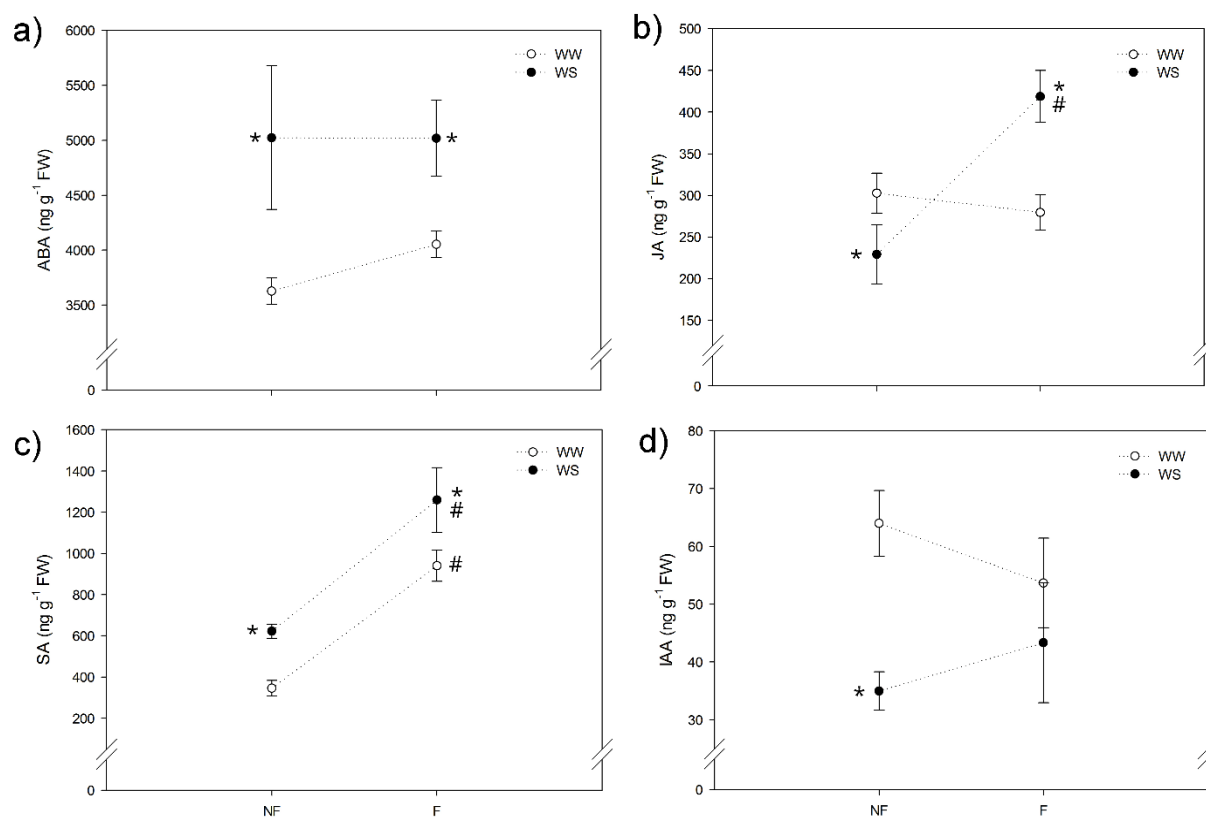


Figure 7 ABA (a), JA (b), SA (c) and IAA (d) content of WW-NF, WS-NF, WW-F and WS-F *E. globulus*. Data are presented as mean ± SE. (*) indicate significant differences within water regimes and (#) indicate significant differences within fungi treatments. ($p \leq 0.05$). WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated

Multivariate approach: a global overview of physiological status

Principal components analysis ordination (PCA) (fig. 8) provides a holistic overview of the experimental setup, revealing a separation between water regimes and fungi treatment. Well-watered plants (WW-NF and WW-F) are mostly grouped together (upper left section) suggesting homogeneity in the biochemistry and physiology response due to higher water potential, gas-exchange parameters (A, gs, E), height and IAA. Water-stressed plants are located in the right biplot section, with non-inoculated plants on the inferior half, due to high proline and ABA content, and fungi-inoculated plants on the upper section due to higher MDA, TSS, SA and JA content.

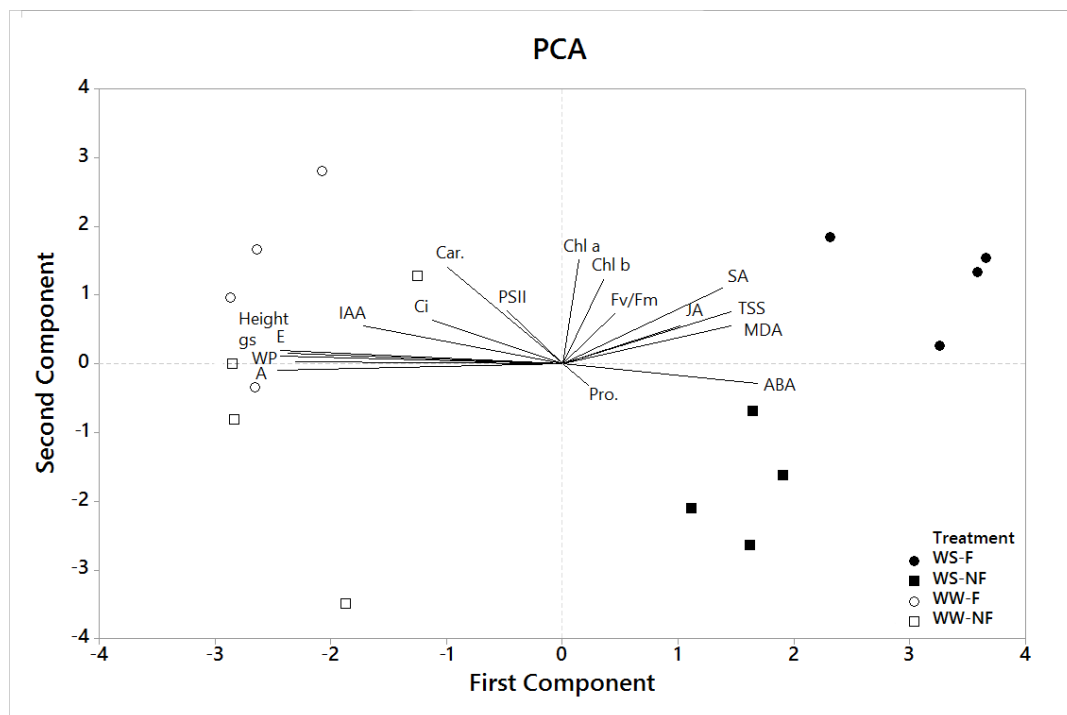


Figure 8 PCA biplot of the physiological data of *E. globulus* during the experimental setup. Different colors are used to differentiate well-watered (white) from water-stressed (black) plants while different symbols are used to differentiate non-inoculated (square) from fungi-inoculated (circle) plants. WW-NF: Well-watered, non-inoculated; WS-NF: Water-stressed, non-inoculated; WW-F: Well-watered, fungi-inoculated; WS-F: Water-stressed, fungi inoculated. Circles are used to highlight the position of specific scores (WW plants; WS-NF and WS-F plants)

Table 3. Two-way ANOVA summary table for morphological and physiological traits of WW-NF, WS-NF, WW-F and WS-F E. globulus. Degrees of freedom (d.f.) and F value are shown for each source of variation; variance (MS) of the residual is also shown. * $P \leq 0.05$, ** $P \leq 0.01$, *** $P \leq 0.001$, NS, non-significant

Parameter	Source of Variation (Sv)	df (Sv, Residual)	F	Significance		
Height (ln)	Water Regime	1,43	236.608	<0.001	***	
	Fungi	1,43	0.0373	0.848	NS	
	Treatment					
	Interaction	1,43	0.0359	0.851	NS	
	MS = 0.00884					
Water Potential	Water Regime	1,21	153.441	<0.001	***	
	Fungi	1,21	0.532	0.474	NS	
	Treatment					
	Interaction	1,21	0.715	0.407	NS	
	MS = 0.00802					
TSS Content	Water Regime	1,13	13.188	0.003	*	
	Fungi	1,13	39.056	<0.001	***	
	Treatment					
	Interaction	1,13	6.449	0.025	*	
	MS = 8.204					
Proline Content	Water Regime	1,9	1.089	0.324	NS	
	Fungi	1,9	0.640	0.444	NS	
	Treatment					
	Interaction	1,9	10.620	0.010	**	
	MS = 8.837					
Gas-Exchange	Ci	Water Regime	1,22	8.870	0.007	**
		Fungi	1,22	3.852	0.062	NS
		Treatment				
		Interaction	1,22	4.860	0.038	*
		MS = 102.620				
	E	Water Regime	1,22	129.900	<0.001	***
		Fungi	1,22	1.216	0.282	NS
		Treatment				
		Interaction	1,22	14.627	<0.001	***
		MS = 0.0194				
	gs	Water Regime	1,22	129.664	<0.001	***
		Fungi	1,22	1.257	0.274	NS
		Treatment				
		Interaction	1,22	9.778	0.005	**
		MS = 0.000192				
	A	Water Regime	1,22	90.177	<0.001	***
		Fungi	1,22	0.0908	0.766	NS
		Treatment				
		Interaction	1,22	12.795	0.002	**

MS = 0.824						
Fluorescence	PSII	Water Regime	1,16	0.583	0.456	NS
		Fungi	1,16	9.132	0.008	*
		Treatment				
		Interaction	1,16	7.561	0.014	*
	MS = 0.0000455					
	Fv/Fm	Water Regime	1,22	0.611	0.443	NS
		Fungi	1,22	0.0862	0.772	NS
		Treatment				
		Interaction	1,22	0.234	0.634	NS
	MS = 0.0000338					
Pigment Content	Chl a	Water Regime	1,12	4.918	0.047	*
		Fungi	1,12	5.048	0.044	*
		Treatment				
		Interaction	1,12	0.00165	0.968	NS
	MS = 0.0233					
	Chl b	Water Regime	1,15	0.402	0.536	NS
		Fungi	1,15	3.787	0.071	NS
		Treatment				
		Interaction	1,15	0.0420	0.840	NS
	MS = 0.00645					
	Carotenoids	Water Regime	1,16	1.457	0.245	NS
		Fungi	1,16	2.342	0.145	NS
		Treatment				
		Interaction	1,16	0.158	0.696	NS
	MS = 0.0103					
MDA Content		Water Regime	1,12	4.359	0.059	NS
		Fungi	1,12	0.419	0.530	NS
		Treatment				
		Interaction	1,12	12.644	0.004	**
MS = 763.14						
Hormones Content	ABA (1/x)	Water Regime	1,11	14.971	0.003	*
		Fungi	1,11	1.478	0.600	NS
		Treatment				
		Interaction	1,11	0.560	0.592	NS
	MS = 0.00000000163					
	SA (ln)	Water Regime	1,12	22.314	<0.001	***
		Fungi	1,12	81.361	<0.001	***
		Treatment				
		Interaction	1,12	2.736	0.124	NS
	MS = 0.0344					
	IAA	Water Regime	1,9	8.380	0.018	*
		Fungi	1,9	0.0217	0.886	NS
		Treatment				
		Interaction	1,9	1.887	0.203	NS
MS = 147.704						

JA	Water Regime	1,12	1.332	0.271	NS
	Fungi	1,12	8.514	0.013	*
	Treatment Interaction	1,12	13.942	0.003	**
			MS = 3246.438		

Discussion

Plants are often simultaneously exposed to multiple stress factors, both biotic and abiotic with different time framings, and, although available literature regarding combinatorial stress is limited, it is suggestive of non-additive interactions with signaling mechanisms appearing to be partially shared (Kissoudis et al., 2014; Walley et al., 2007). Exposure to biotic and abiotic stress leads to a disruption of the plant's metabolism, with physiological costs and, ultimately, to reductions in plant fitness and biomass productivity (Rejeb et al., 2014). The objective of this work was to assess the interaction between a biotic agent (*N. kwambonambiense*), a susceptible host (*E. globulus*) and a favorable environment for disease expression (climate change-driven drought) from a predisposition point of view. Our results conclusively reported that drought lead to predisposition to *N. kwambonambiense* infection in *E. globulus* with increased disease expression and severity.

Plant biomass reduction and growth impairment, perceptible through height, is one of the most noticeable and sensitive variables to be affected by water stress (Chaves and Oliveira, 2004; Shao et al., 2009). Plant growth rates were significantly decreased by water shortage, which is in agreement with other *E. globulus* reports (Correia et al., 2014b; Granda et al., 2011; McKiernan et al., 2014) while WW plants, regardless of *N. kwambonambiense* presence, kept consistently superior heights and corresponding IAA content, which is confirmed by the PCA analysis. Accordingly, water shortage lead to a decrease in midday water potential, as previously described in *E. globulus* (Correia et al., 2014b; Costa et al., 2004; Pita et al., 2003). Although plant water content may have a direct effect on fungal growth, most fungal pathogens are capable to adapt to low water potentials, well below the minimum for growth of most higher plants (Desprez-Loustau et al., 2006; Flowers et al., 2001). While no significant effects on water status due to *N. kwambonambiense* have been found, water-stress may lead to woody tissue desiccation that can result in bark cracks that facilitate pathogen invasion, predisposing the plant to infection while increasing symptom development (Bostock et al., 2014). Consistent with these results, is the extensive window of wound susceptibility found in stressed trees and not in well-watered ones (Mcintyre et al., 1996). Taking this into account, the larger external lesion extensions and higher fungi progression detected on water-

stressed plants can be considered the end result of an increased susceptibility to *N. kwambonambiense* caused by water shortage, also found in the interaction between *Botryosphaeria ribis* and *Eucalyptus* spp. (Old et al., 1990) and in other plant-pathogen interactions (Blodgett et al., 1997; Luque et al., 2002; Van Niekerk et al., 2011).

Water potential decrease leads to a ABA content upsurge, as seen in Correia et al., 2014b and Jesus et al., 2015, an important phytohormone which is transported to the shoot where it induces stomatal closure with costs to the photosynthetic apparatus (Jiang and Hartung, 2008). Stomatal closure leads to a decrease of the CO₂ availability (Chaves et al., 2003), that, together with oxidative stress (Jaleel et al., 2009) impair photosynthesis (Chaves et al., 2009; Lawlor, 2009). Simultaneously, a decrease of the CO₂ mesophyll conductance and transpiration takes place (Flexas et al., 2004). Our results demonstrate this interplay and report an ABA content increase, which is observed in similar *E. globulus* studies (Correia et al., 2014a; Granda et al., 2011) regardless of fungi treatment. Stomatal closure leads to an overall ci, gs, E and A reduction, being in agreement with Guarnaschelli, Prystupa, and Lemcoff 2006 and other reports (Bogeat-Triboulot et al., 2007; Correia et al., 2014b; Warren et al., 2011). Pathogen infection, under WS conditions, leads to a decrease in photosynthetic assimilate production (Berger et al., 2007; Bilgin et al., 2010), which is in agreement with other plant-pathogen interactions (Fontaine et al., 2016; Mayek-Pérez et al., 2002; Scharte et al., 2005; Swarbrick et al., 2006). The subsequent decrease in energy production results in a decay in enzymes and proteins synthesis that play an essential role in pathogen defense, increasing plant predisposition and exacerbating symptom severity (Van Niekerk et al., 2011). On the other hand, gas-exchange parameters increased on WW plants, along with effective quantum yield, with that being suggestive of an overall energy production increase to assure the plant protection. Photosynthetic rate is positively related with chlorophyll content (Anjum et al., 2011) and, although drought is known to lead to a chlorophyll content (Reddy et al., 2004), no content changes were found nor PSII photochemistry was negatively affected in response to water stress or *N. kwambonambiense* infection. Such may be the result of a series of compensatory reactions that mitigate the effects of water stress (Susiluoto and Berninger, 2007) protecting these pigments from degradation, while maintaining or even increasing its concentration as seen in several studies (Correia et al., 2014b; Michelozzi et al., 1995). Such depends upon the role of carotenoids actions as photo and osmoprotectores (Cazzonelli, 2011) and by soluble sugars overall osmotic adjustment mechanisms (Santakumari and Berkowitz, 1991). Furthermore, it reflects the plants ability to adapt to the established conditions, highlighting the need of long-term studies that take into account the acclimation effect (Zhou et al., 2016).

Soluble sugars, besides being involved in osmotic adjustment, have a crucial function as signaling molecules against pathogens through a “high-sugar resistance” mechanism (Morkunas and Ratajczak, 2014) and, upon pathogen infection, an increase of invertase activity takes place to meet such carbohydrate requirement (Roitsch and González, 2004). Soluble sugars are also involved in, or related to, ROS-producing metabolic pathways, protecting plants against photo-oxidation (Couée et al., 2006) whose content early increases upon pathogen infection to give rise to the oxidative burst (Heller and Tudzynski, 2011). Taking both mechanisms into account, the TSS increase upon *N. kwambonambiense* inoculation plants may be considered a direct response to pathogen infection, which is in agreement with other reports (Scharte et al., 2005; Swarbrick et al., 2006) being boosted as a result of water-stress (Couée et al., 2006; Rosa et al., 2009), and an indirect response to the supposed increase of ROS (Fontaine et al., 2016), observed through the high MDA content in the same water-stressed, *N. kwambonambiense*-inoculated plants. This is confirmed by the PCA that conclusively shows higher MDA content in water-stressed, *N. kwambonambiense*-inoculated plants. Nevertheless, water-stressed plants may be unable to properly respond to the harmful oxidative damages triggered by ROS.

Reactive oxygen species are also known to promote SA accumulation, therefore the SA content increase in WS-F was expected (Bostock et al., 2014; Naidoo et al., 2014) and corroborates its role regarding water stress (Hayat et al., 2010), as seen in *Phillyrea angustifolia* and barley (Bandurska, 2005; Munné-Bosch and Peñuelas, 2003) and pathogen local and systemic defense response (Denancé et al., 2013; Halim et al., 2006) as seen in *E. grandis* (Naidoo and Za, 2013). The increase in SA upon *N. kwambonambiense* inoculation in WW conditions may also be preventing an increase of JA in the same conditions, which supports the antagonistic crosstalk (Glazebrook, 2005) similarly detected in *E. grandis* (Naidoo and Za, 2013). Proline accumulation is also involved the response to water stress, thought osmotic adjustment, pathogen infection and overall free radicals scavenging (Chen and Jiang, 2010; Sperdouli and Moustakas, 2012; Szabados and Savoure, 2010). Therefore, the increase in proline upon water stress was expected and is in agreement with other reports (Cha-um and Kirdmanee, 2010; Li and Liu, 2004; Shvaleva et al., 2006), as is the increase in WW plants under *N. kwambonambiense* inoculation, which may reflect an active anti-oxidant response, as seen in other plant-pathogen interactions (Fabro et al., 2004; Haudecoeur et al., 2009). ABA may also have an antagonistic effect over JA (Denancé et al., 2013; Rejeb et al., 2014), which explains the decreased JA content in non-inoculated plants upon WS and is in agreement with other reports with *Eucalyptus globulus* (Correia et al., 2014a) and other plant-pathogen interactions (Anderson et al., 2004; Sánchez-Vallet et al., 2012). Furthermore, JA was found

to play a role in necrotrophic pathogen defense (Glazebrook, 2005; Halim et al., 2006), stimulating induced systemic resistance (Naidoo et al., 2014; Pozo et al., 2008), which explains its increase under WS conditions upon *N. kwambonambiense* inoculation, and is in agreement with other reports (Halim et al., 2004). It also suggests an hemibiotrophic lifestyle, with *N. kwambonambiense* switching from a biotrophic to a necrotrophic interaction, as seen in *Diplodia corticola* (Oliveira Fernandes, 2015) and further corroborates the higher tissue damages in WS plants as living tissue progressively degenerates.

Conclusions

Our results demonstrate that water stress-driven drought caused a decrease in the *E. globulus* ability to respond to *N. kwambonambiense* infection through a predisposition mechanism. Well-watered plants, despite being infected with *N. kwambonambiense*, presented an overall photosynthetic increase (A, PSII), which is suggestive of an active defensive response, through a presumable amplified production of enzymes, proteins and sugars (TSS) required to pathogen defense. *E. globulus* deployed an oxidative (MDA) and high-sugar response (TSS) to prevent pathogen expansion, which is observable through the inferior external and internal fungi expansion parameters. Despite that, oxidative damages were prevented through an active anti-oxidant machinery (TSS, proline) to protect the plant against damages. A concomitant increase of pathogen defense hormones content occurred, particularly salicylic acid, to lead to an efficient defense response, inferior fungi progression and consequent high-growth maintenance.

Water stress lead to a water-potential decrease, which triggered a ABA and TSS induced stomatal closure and overall photosynthetic efficiency decline. Although oxidative (observable through MDA) and high-sugars responses (TSS) took place, the plant was unable to control the resulting oxidative damages. Water potential decrease also induced tissue desiccation that, through a longer window of wound susceptibility lead to a concomitant larger external lesion extension and higher fungi progression. Moreover, and for the first time, under water-stress conditions, *N. kwambonambiense* was found to promote a JA content increase, typical of necrotrophic pathogens, which may suggest an hemibiotrophic lifestyle but also a change in the plant cells status.

This work fills a scientific gap regarding the effect that water stress, a phenomenon that is expected to increase as a result of climate change, on the predisposition of *E. globulus*, a tree established worldwide, with a pronounced economic value, to *N. kwambonambiense*, an opportunist endophytic highly aggressive phytopathogen. Furthermore, this works highlights

the need of long-term water stress studies that take into account the adaptation effects that take place in long-lived plants. Water stress recovery studies should take place to explore if the pathogen returns to a non-aggressive endophytic state, once water status is restored. This work provides insights about the *E. globulus* physiological and hormonal changes, unveiling plant fitness and its role on water stress predisposition that can be of use to the forestry industry.

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Part III

Final Considerations

Drought-disease interactions are of particular relevance to forest trees. Drought-induced diseases are frequently caused by latent endophytic pathogens that hold the ability to be present on the plant's tissue and develop disease only when the host is subjected to stress. This way, typically benign organisms become pathogenic, and water stress becomes a revealer of the pathogen's presence. This interaction and predisposition effect represents a major threat to productive and economically important species. This study is included in that framework and was born from the need to fill a knowledge gap; that is, the effect of water stress on the predisposition of *E. globulus*, a worldwide established tree, with a pronounced economic value, to *N. kwambonambiense*, an opportunist endophytic highly aggressive phytopathogen. An interaction projected to escalate as a result of a climate-change-driven temperature increase and summer precipitation decrease.

The secondary objectives of this work were related with the assessment of the individual effects of water-stress and pathogen infection on *E. globulus*. The main objective was to assess the water stress predisposing effect on *E. globulus* and find out if that would result in an increased susceptibility to *N. kwambonambiense* infection and disease development. For that, all plants were firstly subjected to an acclimation period. After that, two water regimes were established, a well-watered and a water-stressed one. Once water-stressed plants achieved 18% of FC, one well-watered and one water-stressed groups were inoculated with *N. kwambonambiense*. At the end of the experiment, several morpho-physiological and biochemical parameters were assessed.

After measuring growth, external lesion extension, water potential, lipid peroxidation, osmolyte and hormone quantification, we were able to conclude that water stress was the main contributor to growth impairment and, therefore, water-stress associated parameters' changes. Well-watered plants (Fig. 1a) presented superior heights, consistent with the IAA content. Despite being infected with *N. kwambonambiense*, presented an overall photosynthetic-parameters' improvement (A, gs, E, Φ PSII). Such allowed plant protection through the production of sugars (TSS), proline and hormones (salicylic acid). Oxidative damages, (partially observed through MDA content), to have happened, were avoided, in part, due to the action of proline and soluble sugars. Such allowed a high-growth maintenance and inferior fungi internal and external progression.

Water-stress (Fig. 1b) decreased water potential, which together with an ABA content increase, lead to stomatal closure. Plants, although experiencing an increase of intercellular

CO₂ concentration, were found unable to use it with a consequent photosynthetic decline (A, E, gs). Such lead to a presumable decrease in the production of enzymes and proteins required in pathogen defense mechanisms and therefore, impairment of the plant's ability to properly react to fungi infection. Oxidative damages (partially observed through MDA content) weren't properly managed and further affected *E. globulus*. A SA content increase was found and, for the first time, under water-stress conditions, *N. kwambonambiense* was found to promote a JA content increase, typical of necrotrophic pathogens. Such may suggest a lifestyle change from biotrophic to necrotrophic as plants cells progressively degenerate.

It is possible to conclude that water stress negatively influenced *E. globulus*, affecting its ability to respond to *N. kwambonambiense* infection. Such is justified by the consistently larger external lesion extensions found in water-stressed, *N. kwambonambiense*-inoculated plants. An increasing number of studies have been revealing the diversity and functional importance of fungal endophytes in trees. This study further represents a clear example of a positive association between drought and disease, i.e., drought or drought and disease acting synergistically affecting tree health. A matter of concern that should be further studied as the effects of climate-change are becoming more severe.

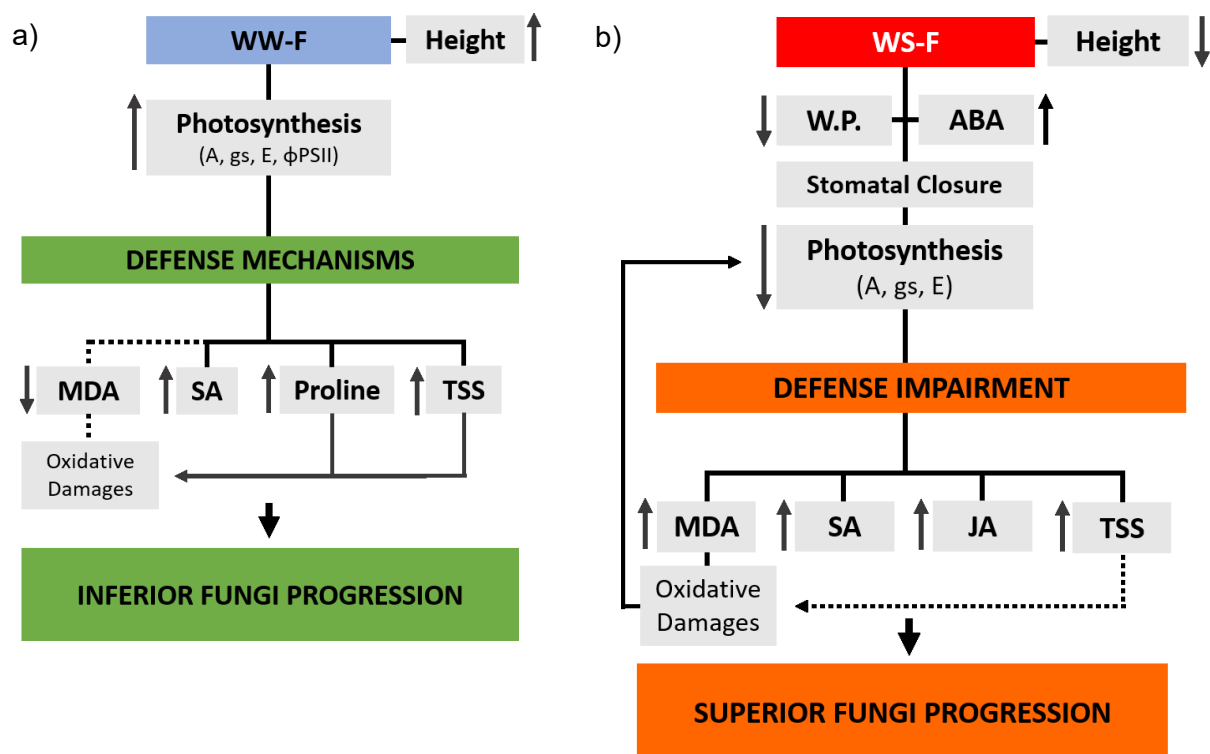


Figure 1 Graphical representation of the obtained results of well-watered and water-stressed *N. kwambonambiense*-inoculated plants

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